#### STIC-Biotech/ChemLib

From: Sent: To:

Subject:

McGarry, Sean Tuesday, June 17, 2003 10:12 AM STIC-Biotech/ChemLib SEQ SEARCH 09/964,666

JUN 17 2003

(STIC)

Sean McGarry 73484 AU 1635 CM1 11D07 Office CM1 11E12 Mailbox 305-7028

09/964,666

Please, a length limited (nt ≤ 50) search of SEQ ID NOS 9-14.

Thank You

Edward Hart Technicai Info. Specialist STIC/Biotech CMI 6B02 Tel: 305-9203

Searcher:
Phone:
Location: / MM2
Date Picked Up:
Date Completed: 15/23/603
Searcher Prep/Review:
Clerical:
Online time:

TYPE OF SEARCH:
NA Sequences:
AA Sequences:
Structures:
Bibliographic:
Litigation:
Full text:
Patent Family:
Other

VENDOR/COST (where ap STN:	plic.)
DIALOG:	
Questel/Orbit:	
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June

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2003, 21:23:11; Search time 522.031 Seconds (without alignments) 1449.478 Million cell updates/sec

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nucleic search, using sw model

Copyright

GenCore version 5.1.6 (c) 1993 - 2003 Compugen Ltd

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Minimum

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Searched:

Total number

of hits satisfying chosen parameters:

841850

2054640 segs, 14551402878 residues

Scoring table: Sequence: Perfect score: Title:

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US-09-964-666-10 26

TGGTGCATGTCTTTGGTCCCAGCTAC

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Database

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score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.

SUMMARIES

#### RESULT 1 AX050293 LOCUS DEFINITION ACCESSION SOURCE ORGANISM VERSION KEYWORDS REFERENCE Result 000 000 0000 O 000 0000 JOURNAL TITLE AUTHORS Ö 15. 6 15. 6 15. 15. Score 14.8 14.8 14.8 16 AX050293 Sequence 47 AX050293 AX050293.1 Shimkets,R.A., Fernandes,E. and Boldog,F. Secreted polypeptides and corresponding polynucleotides Patent: WO 0070046-A 47 23-NOV-2000; Curagen Corporation (US) synthetic construct. synthetic construct artificial sequences. 1 (bases 1 to 21) Match Query 47 Length 21 bp from Patent WO0070046. GI:12226574 DΒ AR078070 E35651 AR078068 AR098737 E35649 AX010420 AR022354 AX248371 AX249312 AX405134 AX477864 HSA224254 AX184141 AX116084 AR164904 AR116815 AR067246 AX117706 AX429427 AX050293 AX199678 AX378267 AX476157 AX476158 I89866 AX291119 AX249648 AX040679 AX173376 I15450 I41297 I31358 AR040575 AR055599 BD002550 AX234354 AR169512 AX476156 E35696 AR171891 AR023949 AR040576 ALIGNMENTS DNA linear AR023949 Sequence 112127 Sequence 32 115450 Sequence 28 141297 Sequence 32 131358 Sequence 27 132222 Sequence 18 AX010420 AR023354 AX2483312 AX249312 AX405134 AX4071334 AX40679 AX184141 AX040679 AX173376 AX291119 AX249648 AX117706 AX429427 AX116084 AR164904 BD002550 AX234354 AR169512 AX476156 E35696 De AR171891 AR078070 E35651 De Description AR055599 AR040575 AX476158 AX050293 PAT 12-JAN-2001 158 Sequence 5 Sequence 24 550 Gene comp 070 Sequence 1 Detection o 068 Sequence 737 Sequence Detection o Sequence Sequence

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                                 Cohen,D., Blumenfeld,M., Chumakov,I., Abderrahim,H. Obesity associated biallelic marker maps Patent: WO 0206525-A 56 24-JAN-2002;
                                                                                                                                                  AX378267 47 bp
Sequence 56 from Patent WO0206525
AX378267
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Nucleic acids containing single nucle
methods of use thereof
Patent: WO 0151670-A 608 19-JUL-2001;
Curagen Corporation (US)
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Mammalia; Eutheria;
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/db_xref="taxon:32630"
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1 (bases 1 to 21)
Picoult-Newburg,L. and Pohl,M.
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Evans.G.A. and Smith,M.W.
Method for generation of sequence sampled
Patent: US 5851760-A 594 22-DEC-1998;
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Patent: WO 0129262-A 1207 26-APR-2001;
Orchid BioSciences, Inc. (US)
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Picoult-Newburg, L. and Pohl, M.
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105 from patent US 6274339.
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/db_xref="taxon:32630"
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Pred. No. 4.4e+04;
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Methods and compositions for the diagnosis
weight disorders, including obesity
Patent: US 6274339-A 105 14-AUG-2001;
                                                                                           Unknown
                                                                                                                                                                                                                                                   18; Conserv
                                                                                                                                                                                                                                                                                                                                              Unclassified.

1 (bases 1 to 44)

Burgess, R., Friedrich, G., Zambrowicz, B.
Targeted gene discovery
Patent: US 6139833-A 10 31-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                  Unknown
                                        5' to 3' exonuclease mutations of thermostable Patent: US 5795762-A 28 18-AUG-1998;
                                                   1 (bases 1 to 36)
Abramson, R.D. and Gelfand, D.H.
5' to 3' exonuclease mutations
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PAT 16-MAY-2001

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5' to 3' exonuclease mutations of thermostable
Patent: US 5466591-A 28 14-NOV-1995;
Location/Qualifiers
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Gelfand,D.H., Lawyer,F.C. and Stoffel,S.
Mutated thermostable nucleic acid polymerase
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Location/Qualifiers
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Pred. No. 5.3e+04;
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                 PAT 13-MAY-1997
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I41297
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Gelfand,D.H., Lawyer,F.C. and Stoffel,S.
Purified thermostable nucleic acid polymerase
              CTP extended form of glycoprotein ho Patent: US 5585345-A 18 17-DEC-1996; Location/Qualifiers 1. 41
                                                                                     Unknown
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Patent: US 5582979-A 270 10-DEC-1996;
Location/Qualifiers
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19; Conserv
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BASE COUNT 4 a ORIGIN  Query Match Best Local Similarity Matches 18; Conserv	/organism="unknown" 4 a 12 c 14 g 11 t 55.4%; Score 14.4; DB 6; Length 41; milarity 75.0%; Pred. NO. 8e+04; Conservative 0; Mismatches 6; Indels 0; Gaps 0;
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Search completed: June 21, Job time : 527.031 secs	d: June 21, 2003, 22:14:28 031 secs
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## ALIGNMENTS

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or genetic mapping of phenotypic traits	phenotype, in forensics, paterni	New nucleic acid segments containing polymorphic sites - used for,		WPI; 1999-080963/07.		Berno A, Chee M, Fan J, Lipshutz RJ;		(AFFY-) AFFYMETRIX INC.	24 OON 1997, 9703 0030394.		22-JUN-1998; 98WO-US12930.		30-DEC-1998.		WO9858529-A2.		Homo sapiens.			Polymorphism; biallelic; paternity testing; forensic; genetic mappin		Human biallelic polymorphic DNA fragment WI-17197.	31-MAR-1999 (first entry)		AAX06149;	AAX06149/c AAX06149 standard; DNA; 31 BP.	ULT 1

Claim 1; Page 11; 61pp; English

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RESULT
ABL4096
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      application programme and including records, each record comprising information identifying a polymorphism shown in the above sequences. The products and methods can be used for analysing polymorphic sites in individuals for testing for the presence of a disease phenotype or in forensics, paternity testing or genetic mapping of phenotypic traits. They can also be used for the production of polypeptides expressed by variant genes and for the production of transgenic animals. The nucleic acid segments can also be used in the manufacture of medicaments for the
The invention relates to a novel human transcription regulation factor ZFM1 isomer 19.47 and polynucleotides encoding the protein. The protein can be expressed by standard recombinant methodology. The protein and
                                                                                                                                                   Human transcription regulation factor ZFM1 isomer 19.47 and encoding polynucleotide, used in diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and inflammation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Transcription regulation factor ZFM1 isomer 19.47; human; ZFM1; transcription regulation factor; cytostatic; haemostatic; virucide; immunomodulatory; antiinflammatory; gene therapy; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 medium for storing data for access by an application programme being executed on a data processing system. Such a method comprises a data structure stored in the computer-readable storage medium, the data
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequences AAX06101-X06558 represent human DNA fragments which cont biallelic polymorphic markers. The base occupying the polymorphic is indicated by the appropriate IUPAC-IUB ambiguity code. These fragments can be used in a method for determining polymorphic form an individual. The invention further provides communications.
                                                                                                   Example 4; Page 21; 38pp; Chinese
                                                                                                                                                                                                                                                                             WPI; 2002-258025/30
                                                                                                                                                                                                                                                                                                                                                                                                                             28-JUN-2000; 2000CN-0116826
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-JUN-2001; 2001WO-CN01050
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                                                                                                                                                                                                                                                                                                                                                                             (SHAN-) SHANGHAI BIOWINDOW GENE DEV INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         treatment or
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76.9%;
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Best Local
                                                          pharmacogenomics. Specifically, the SECX polypeptides and nucleic acids are useful for treating cancer; other disorders related to angiogenesis e.g. abnormal wound healing, psoriasis; neurodegenerative diseases; immune disorders; liver cirrhosis; benign tumours; fibrocystic conditions and tissue hypertrophy (e.g. benign prostatic hypertrophy); allergic reactions and conditions such as asthma and other respiratory problems; as well as in treating or preventing diseases associated with organ
                                                                                                                                                                                                    primers and probes used in the isolation and characterisation of DNA sequences of the invention. The new polypeptides and nucleic can be used in screening assays, detection assays, preventive or
                                                                                                                                                                                 can be used in screening assays, detection assays, preventive or predictive medicine, therapeutic and prophylactic treatment, and
                                                                                                                                                                                                                                                                                                                                                         New SECX polypeptides and nucleic acids useful for treating or preventing cancer, other disorders related to angiogenesis, neurodegenerative diseases, autoimmune disorders and allergic reactions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                encoding polynucleotides are used in diagnosis and treatment of malignant tumours, haemopathy, HIV infection, immunological diseases and inflammation. The present sequence represents the human transcription regulation factor ZFM1 isomer 19.47 cDNA cloning PCR primer.
                            polypeptides can also be used for bone, cartilage, tendon,
                                                                                                                                                                                                                                                      proteins
                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-025020/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SECX; secreted protein; cancer; angiogenesis; wound healing
                                               transplantation, atherosclerosis-associated diseases or disorders.
                                                                                                                                                                                                                                                    Polynucleotide sequences AAF23410 - AAF23419 encode secreted SECX proteins AAB49649 - AAB49658. Sequences AAF23420 - AAF23450 represent
                                                                                                                                                                                                                                                                                                        Example 8; Page 117; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-MAY-2000; 2000WO-US13291
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     respiratory
PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Forward PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-MAR-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-JAN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       immune disorder;
              tissue growth or regeneration,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 œ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ഗ
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19; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   rder; neurodegenerative disease; allergic reaction;
problem; organ transplantation; contraceptive; human;
proliferative disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99US-0134315.
2000US-0175744.
2000US-0188274.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primer for amplification of DNA encoding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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86.4%;
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Pred. No. 1
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            tissue repair and
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RESULT 4
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Best Local 9
                                      Query Match
Best Local (
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                            Matches
                                                                                                                                      Alzheimer's disease, Down's syndrome, oeuema, rapro compania, arthritis, glomerulonephritis, haemolytic anaemia, thrombocytopaenia, arthritis, meningitis, muscular disorders, dementia, neurological diseases, tube meningitis, muscular disorders, dementia, blood pressure disorders
                                                                                                                                                                        The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of diseases associated will aberrant protein expression, including cancer, amyloidosis, diabetes, Alzheimer's disease, Down's syndrome, oedema, lupus (SLE), vasculitis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   inhibit thromboses, infections caused by bacteria, virus, fungi and parasites, and as a vaccine. SECX antibodies may be used to isolate detect SECX proteans, monitor protein level in tissue as part of a clinical testing procedure, treat proliferative disorders including
                                                                                                  sclerosis, male infertility, hypercalcaemia, blood pressure disorders, osteoporosis, pathogenic infections, hypercholesterolaemia, obesity or autoimmunity. The present sequence is a polymorphism-containing oligonucleotide fragment of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                        07-JAN-2000; 2000US-0174962
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                                                                            Sequence
                                                                                                                                                                                                                                                          Claim
                                                                                                                                                                                                                                                                                         Isolated human polynucleotides containing single nucleotide polymorphisms, useful for the treatment and diagnosis of e.g. \,
                                                                                                                                                                                                                                                                                                                                  P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                       Shimkets RA, Leach MD;
                                                                                                                                                                                                                                                                                                                                                                                               (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                               05-JAN-2001; 2001WO-US00322.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         tumours and benign hyperplasias.
                                                                                                                                                                                                                                                                                                                                              2001-451871/48
                           '19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               single
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18; Conser
3 GTGCATGTCTTTGGTCCCAGCTAC
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                           Conservative
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85.7%;
                                      61.5%;
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Pred. No. 4.8e+02;
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ies 5;
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RESULT 6
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a set of novel map-related biallelic markers, preferably located on obesity disorder-associated chromosomal regions on chromosomes 3, 10 and 19. The markers are useful for genotyping or estimating the frequency of an allele in a population, for detecting an association between a genotype or haplotype and a phenotype, e.g. a disease involving drug responses, obesity or disorders related to obesity, such as hyperuricaemia, digestive pathology, hepatic function disorders, cancer, cardiovascular disease, hypertension, hyperlipidaemia, insulin disorders, atheromatous disease and cardiac insufficiency. The marker allele and a phenotype and/or between a biallelic marker allele and a phenotype and/or between a biallelic marker allele and a phenotype and/or between a biallelic marker baplotype and a phenotype. This sequence represents a human obesity-associated biallelic marker located on chromosome 10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Set of novel map-related biallelic markers, preferably located obesity disorder-associated chromosomal regions on chromosomes and 19, useful, for e.g. detecting statistical correlations be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI;
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                                                                              AAQ82594;
                                                                                                                                                                 AAQ82594 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 47
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1e+03;
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14-SEP-1995

(first entry)

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RESULT 7
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Best Local
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                                                                                   Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease;
                                                          acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis;
                                                                                                                                                                                                                                                                                                        14-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                      AAH40033
                                                                                                                                                                                                                                                                                                                                                                                                                              AAH40033 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      high density sequence-based maps, and ultimately, for the complete sequencing of genomic DNA directly from cosmid clones. See AAQ82001-Q82706 and AAQ91325-Q91358 for STS primers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequencing
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07-SEP-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    encing complex genomes, present as fragments in a cosmid ary - by sequencing end-specific nucleotides of each clone correlating with spatial relationship of cosmid, esp. for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20
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17; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATGTCTTTGGTCCCAGCTAC 26
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ibrary; chromosome
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93US-0117952.
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                                                                                                                                                                                                                                              PCR primer SEQ ID
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85.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 C; 9 G;
                           investigation; paternity analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               genomic analysis; complex genome mapping;
se 11; sequence tagged site; STS analysis;
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                        PCR primer; ss
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RESULT 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, familial hypercholesterolaemia, polycystic kidney disease, osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial disease of which a component is or may be genetic such as autoimmune diseases, including, rheumatoid arthritis, multiple sclerosis, inflammation, cancer, nervous system diseases and infection by pathogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The
                                                                          Genomic analysis;
                                                                                                          Genomic
                                                                                                                                                                                                          ABK92971 standard; DNA; 27
                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21
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                                           Synthetic
                                                                                                                                          22-AUG-2002
                                                                                                                                                                            ABK92971
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     microorganism. The method is also useful in forensic investigations and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             oligonucleotides are useful for determining the presence, absence or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 64; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New genotyping oligonucleotide, useful for detecting the presence,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    paternity analysis. The present sequence represents a PCR
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                                                                                                          method PCR primer 3'DEGIS431
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85.0%;
                                                                          primer;
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                Claim 11; Page 40; 100pp; English
                                                                Preparation of biologically-activated circularly-permuted proteins by scanning permutagenesis for generating libraries of permuteins with improved therapeutic properties -
                                                                                                                                                                                                                             Lee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200018905-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Biologically-activated circularly-permuted protein; permutein library generation; therapeutic property;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA35578
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Permutein linker
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAA35578 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-MAY-2002
                                                                                                                                                                                                                                                                           (SEAR ) SEARLE & CO G D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (MBIF-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      invention describes a method of analysing a DNA sample using atton, digestion and extension techniques. The method is also ful for identification and analysis of unknown sequences upstream a known region or sequence. This sequence represents a PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      in analysis of sequences captured from a st of bacterial genomic DNA.
                                                                                                                                                                           2000-293145/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2002-489951/52
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MBI FERMENTAS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TGGTGCATGTCTTTGGTCCCAGCTA 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TGGTGCATATTCTTGAACCCANCKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page 21; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 6 A; 6 C; 4 G; 8 T; 3 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2000US-0695545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2001WO-CA01487
                                                                                                                                                                                                                                                                                                                              98US-0101908
                                                                                                                                                                                                                                                                                                                                                                                 99WO-US20891
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         iry generation; therapeutic improve bio-distribution; !
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           58.5%;
68.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nucleotide sequence FGS3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 15.2; DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         half life;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                restriction endonuclease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    permutein; linker;
antigen;
ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 γď
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAL31378/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC (permuteins) comprises the use of a method comprising making a series of CC (circularly permuted genes. The circularly permuted genes are inserted into a display vector, where they are expressed so that the proteins CC of display vector where they are expressed so that the proteins CC of display vectors presenting the expressed circularly permuted protein series is generated. A target protein that can bind a biologically active CC circularly permuted protein can be used to affinity select the gresenting display vectors. The selected display vectors can be isolated CC and analysed to identify the presented circularly-permuted protein. The CC permuteins conform to the structure of a parent protein consisting of a compared to the carboxy portion of the parent protein, and CC segment derived from the carboxy portion of the parent protein, and CC segment derived from the amino terminus of the parent protein, and CC segment derived modelinks and analysed to coreate the permuteins with improved therapeutic properties compared to CC create the permuteins with improved therapeutic properties compared to CC libraries of permuteins with improved therapeutic properties compared to CC immunotherapy as probes or intermediates used to construct other useful comparations and becomes a permutein and biological and therapeutic of the construct other useful comparations are not permuteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                  Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antimicrobial; gene therapy; vaccine; amylase; can amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating f complement related protein; cytochrome; kinesin; cytokine; interfer interleukin; G-protein coupled receptor; thioesterase; inflammation multifactorial disease; autoimmune disease; infection;
                                                                                                                                                        28-DEC-1999;
27-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAL31378;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAL31378 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     permuteins. Permutei properties compared
                                                 WPI; 2001-465210/50.
                                                                                                                                                                                                           28-DEC-2000;
                                                                                                                                                                                                                                              05-JUL-2001
                                                                                                                                                                                                                                                                              WO200147944-A2
                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                     nervous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human SNP oligonucleotide #4586.
                                                                                                                   (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-JAN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    rmuteins. Permuteins have improved biological and therapeutic operties compared to their two individual components due to a bio-distribution or half-life.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            6 CATGTCTTTGGTCCCAGCTA 25
                                                                                                                                                                                                                                                                                                                                                   system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CATGTCTTTGGGCCCTGCCA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                      99US-0173419
2000US-0173419
                                                                                                                                                                                                           2000WO-US35498
                                                                                                                                                                                                                                                                                                                                                   disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Α,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                58.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15 C;
                                                                                                                                                                                                                                                                                                                                                   SS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10 G; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15.2;
No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
                                                                                                                                                                                                                                                                                                                                                                                      inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        to alterations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                            cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                         factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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Polymorphic nucleic acids encoding e.g. amylases, cyclins, poncogenes and histones, useful for diagnosing and treating,

cyclins, polymerases,

e.g.

cancer, autoimmune diseases and infections

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RESULT 11
AAH38411/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         proteins, cytochromes, kinesins, cytokines, interferons, interleukins, 6-protein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases with a genetic component, such as autoinmune diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney, leases of the nervous system and an infection of pathogenic
primer extension
                                                                   Claim 1; Page 56; 83pp; English
                                                                                                                    New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nuc
                                                                                                                                                                                                                    Picoult-Newburg L,
                                Sequences AAH37205 -
                                                                                                                                                                                                                                                       (ORCH-) ORCHID BIOSCIENCES INC
                                                                                                                                                                                                                                                                                             15-OCT-1999;
                                                                                                                                                                                                                                                                                                                               13-OCT-2000; 2000WO-US28436
                                                                                                                                                                                                                                                                                                                                                                                                      WO200129262-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Single nucleotide polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SNP specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAH38411 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases, amyloid proteins, angiopoietin, apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related
                                                                                                                                                                                                                                                                                                                                                                    26-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            polycystic kidney disease; osteogenesis imperfecta; autoimmune disease;
acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer;
Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                  2001-290930/30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 10 A; 13 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SNPE primer SEQ ID
                                                                                                                                                                                                                                                                                             99US-0160096
               (SNPE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                      Pohl M;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     58.5%;
             AAH40944 represent IPE) primers, and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           investigation; paternity analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 15.2;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide primer extension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1207
             PCR primers,
e sequences o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.6e+03;
3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
               of
           single nucleotide f regions flanking
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      50;
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                                                                                                                          a nucleic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 12
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Best Local S
Matches 18
                                                                                                                                                                                                                             21-JUL-1998;
20-OCT-1998;
05-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group of individuals, having a pathological phenotype trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial disease of which a component is or may be genetic such as autoimmune diseases, including, rheumatoid arthritis, multiple sclerosis, inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a single nucleotide primer extension (SNPE) primer specific for a human SNP containing DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mahogany gene; mouse; mg gene; regulatory defect; gene therapy; obesity; weight regulation; cell therapy; body weight disorder; cachexia; anorexia; hyperpigmentation; increased metabolic rate disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAZ92014 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscul
dystrophy, familial hypercholesterolaemia, polycystic kidney disease,
                                                                                                                                                                                                                                                                                                                                      03-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mahogany protein gene exon 28+UTR primer CelegE170.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ92014;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 27 BP; 7 A; 7 C; 6 G; 6 T; 1 other;
                                                                                                                                                           Moore
                                                                                                                                                                                            (MILL-) MILLENIUM PHARM INC
                                                                                                                                                                                                                                                                                                   21-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                        WO200005373-A2
                                                                                                                                                                                                                                                                                                                                                                                                             Mammalia
                                                                                                                                                                                                                                                                                                                                                                                                                                            hyperphagia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-JUN-2000
                                                                    creatment
                                                                                                                        2000-195103/17.
                                                                                                                                                         ζ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 TGGTGCATGTCTTTGGTCCCAGCT 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TAGGGCAAGTCNTTAGTCCAAGCT
                                                                  and murine mahogany genes, useful,
of body weight disorders -
                                                                                                                                                         Nagle DL;
                                                                                                                                                                                                                                                                                                                                                                                                                                            hyperpigmentation; increased metabolic rate ia; Antiobesity; antianorexic; anticachexic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                             98US-0093630.
98US-0104978.
99US-0245041.
                                                                                                                                                                                                                                                                                                   99WO-US16484
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      57.7%;
75.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ..8e+03;
les 6;
                                                                                    e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                      for
                                                                                    diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         muscular
                                                                                    and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                              SS
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This sequence represents a PCR primer

for a

mahogany gene of

the

Example;

Fig 5;

188pp;

English

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AAX36190/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 18
                                                                              The specification describes an infectious virus which comprises a recombinant viral genome and a chimeric integrase. The chimeric integrase can provide an infectious recombinant virus that is capable of infecting higher eukaryotic cells and incorporates a chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             levels (or function) of (II) (for diagnosis, prognosis or monitoring of treatment); to evaluate (II)-expressing cells intended for cell therapy, and as therapeutic mg inhibitors. Cells that express the mg gene (or contain the mg polypeptide) are used to identify agents (A) that modulate mg activity. (A) are potentially useful for the treatment of body weight disorders, particularly obesity, cachexia or anorexia, or other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             invention. The mahogany genes are used: (i) to produce recombinant mahogany (mg) proteins (II); (ii) as a source of antisense, ribozyme or triplex-forming therapeutics; (iii) as a source of diagnostic probes and primers for detecting expression of mg genes or mutations, regulatory defects, in this gene, or for isolation of related sequences; and (iv) in (cell-based) gene therapy. (II) are used to raise specific antibodies (Ab); to identify other (extra)cellular products involved in weight regulation, and to screen for agents that disrupt interaction between (II) and other macromolecules. The Ab are used to detect abnormal
                                                                                                                                                                                                                                                       WPI; 1999-287449/24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mutagenesis; gene trapping; retroviral integration;
                   mutagenesis,
                                                                                                                                                                   Disclosure; Fig 10B; 95pp; English.
                                                                                                                                                                                                                                                                                                                                                                                          06-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                           18-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Infectious virus; recombinant viral genome; chimeric integrase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX36190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX36190 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                conditions associated with the mg gene such as hyperpigmentation, hyperphagia and disorders that result in increased metabolic rate
                                 The viruses can be used
                                                      viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9907389-A1
                                             integrase activity that has been engineered to target and guide viral integration to specific regions of the genome of living cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Intronic location
                                                                                                                                                                                                      chimeric
                                                                                                                                                                                                                      infectious viruses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 l Similarity
18; Conser
                                                                                                                                                                                                                                                                                                                        LEXICON GENETICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           formation; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TGGTTCATGCTTTTTATCCCAGC 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TGGTGCATGTCTTTGGTCCCAGC 23
                                                                                                                                                                                                     integrase, used
                                                                                                                                                                                                                                                                                       Friedrich
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
 target sequences
               via gene trapping, of genes encoding,
                                                                                                                                                                                                                                                                                                                                                       97US-0907598
                                                                                                                                                                                                                                                                                                                                                                                          98WO-US16373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        where the LTR trapping construct integrates.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; 5 C; 6 G; 4 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            57.7%;
78.3%;
                                                                                                                                                                                                                                                                                       GA,
             in methods for the rapid identification and rapping, of genes encoding, regulated by, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               44
                                                                                                                                                                                                   comprising a :
d particularly
                                                                                                                                                                                                                                                                                       Sands
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score
Pred.
 for DNA binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                       AT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               No.
                                                                                                                                                                                                   a recombinant viral genome and rly for targeted gene discovery
                                                                                                                                                                                                                                                                                       Zambrowicz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.8e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 21;
proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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RESULT 14
AAL29694
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Best Local :
The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases, amyloid proteins, angiopoiețin, apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related proteins, cytochromes, kinesins, cytokines, interferons, interleukins, G-protein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases with a genetic component, such as autoimune diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes,
                                                                                                                                                                                                                                                          Polymorphic nucleic acids encoding e.g. amylases, oncogenes and histones, useful for diagnosing and cancer, autoimmune diseases and infections -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antimicrobial; gene therap; vaccine; amylase; can amyloid protein; angiopoletin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating f;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 random integration of viral DNA into the host genome is reduced. The allows for a more efficient and high throughput analysis of target sites. The present sequence represents the sequence of the intronic location where the LTR trapping construct integrates.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  for retroviral integration and provirus formation to occur at a location within the genome. By deleting the naturally occurring nonspecific DNA binding domain of the integrase, the amount of
                                                                                                                                                                                                                            Claim 1;
                                                                                                                                                                                                                                                                                                                                                                Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          multifactorial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       complement related protein; cytochrome; kinesin; cytok interleukin; G-protein coupled receptor; thioesterase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 44 BP; 9 A; 14 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                   27-DEC-2000; 2000US-0173419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-DEC-2000; 2000WO-US35498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200147944-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nervous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human SNP oligonucleotide #2902.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAL29694 standard; DNA; 50
                                                                                                                                                                                                                                                                                                                                                                                                (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 TGGTGCATGTCTTTGGTCCCAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                          Page 2217; 4143pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TAGCGCATGCCTTTAGTCCCACC 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first
                                                                                                                                                                                                                                                                                                                                                                Leach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-0173419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  57.7%;
78.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         autoimmune disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12 G; 9 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   other,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    <u>ن</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cytokine;
                                                                                                                                                                                                                                                                            cyclins, polymerases, treating, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       interferon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      at any
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        factor;
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RESULT 15
ABN87593/c
ID ABN87593;
ABN87593;
XX
C ABN87593;
XX
DT 07-AUG-2002 (first entry)
XX
DT 07-AUG-2002 (first entry)
XX
DT 07-AUG-2002 (first entry)
XX
DE Oligodendrocyte-myelin glucop
XX
Human; oligodendrocyte-myelin
XX
ROS Homo sapiens.
XX
SOS HOMO sapiens.
XX
PN CN1333225-A.
XX
PN CN13332
Search completed: June 21, 2003, 22:26:21 Job time: 145:594 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         δõ
                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes human oligodendrocyte-myelin glucoprotein 9.68 (I). Also described is a method for producing (I) using DNA recombination technology. (I) can be used in the treatment of several diseases, such as nervous system dysfunction related diseases. The present sequence represents a probe for (I), which is used in an example from the present invention.
                                                                                                                                                                                                                                                                                                                                                                    Sequence 41 BP; 5 A; 8 C; 16 G; 12 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 6; Page 22 (Disclosure); 34pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel polypeptide--oligodendrocyte-myelin glucoprotein 9.68 and polynucleotide for encoding the polypeptide -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-JUL-2000; 2000CN-0117081.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-JUL-2000; 2000CN-0117081.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; oligodendrocyte-myelin glucoprotein 9.68; probe; ss; nervous system dysfunction related disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oligodendrocyte-myelin glucoprotein 9.68 probe 2 SEQ ID NO:9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 50 BP; 14 A; 11 C; 12 G; 13 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney, leukaemia), diseases of the nervous system and an infection of pathogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (SHAN-) SHANGHAI BIODOOR GENE DEV CO LTD
                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 TGGTGCATGTCTTTGGTCCCAGC 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18; Conserv
                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                      56.9%;
73.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     57.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ВP
                                                                                                                                                                                                                                            0; Mismatches
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                                                                                                                                                                                                                                      Score 14.8; DB 24;
Pred. No. 2.3e+03;
0; Mismatches 7;
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Maximum Match 100%
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1: /cgn2_6/ptodata/l,
2: /cgn2_6/ptodata/l,
3: /cgn2_6/ptodata/l,
4: /cgn2_6/ptodata/l,
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/cgn2_6/ptodata/1/ina/BCTUS_COMB.seq:*
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US-07-971-819A-32
US-07-971-819A-32
US-08-458-813-28
PCT-US91-07035-28
PCT-US91-07035-28
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US-08-222-177A-270
US-08-222-177A-270
US-08-228-952-12
US-09-286-959B-12
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Sequence 11
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67.7 76.9 76.9 ative CTTTGG 	ication US/09641638  N: nfeld, Marta meleret, Lydie kov, Ilya N: Annick N: BIALLELIC MARKE N: GENES INVOLVED GENSET.051CP1 CON NUMBER: US 09/502 1000-02-11 NUMBER: US 60/133 1999-05-07 NUMBER: US 60/133 1999-03-23 1999-03-23 1099-03-23 11999-03-12 NOS: 1304pm apiens apiens apiens apiens apiens apiens continuous (10-500-258 : poleature) eature N: n=a, g, c or t		3 4 4 9 9 5 b	47 50 18	29 29 32	40 32
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re 17.6; d. No. 13; Mismatches TAC 26     TAC 38	IN 1.6 1.6	ALIGNMENTS	3-48 3-905 3-471 3-471 3-178 3-257	0-641 043-3 0-071	19-489 19-302 19-609 18-295 18-948 18-362	-927 -987 -987
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RESULT 2
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; Sequence 594, Application US/08117952
; Patent No. 5851760

ENERAL INFORMATION:

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Best Local Similarity
Matches 17; Conservi
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                                                                      EARLIER APPLICATION NUMBER: 60/104,978 EARLIER FILING DATE: 1998-10-20 NUMBER OF SEQ ID NOS: 131
                                                                                                                     CURRENT APPLICATION NUMBER: US/09/245,041
CURRENT FILING DATE: 1999-02-05
EARLIER APPLICATION NUMBER: 60/093,630
EARLIER FILING DATE: 1998-07-21
                                                                                                                                                                                                 TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR THE DIAGNOSIS AND TREATMENT
TITLE OF INVENTION: OF BODY WEIGHT DISORDERS INCLUDING OBESITY
FILE REFERENCE: 7853-136
                                                                                                                                                                                                                                                                                       ENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                           09-245-041-105/c
TYPE: DNA
                                                         SOFTWARE:
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atent No. 6274339
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SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
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FILING DATE: 15-1091
ATTORNEY/AGENT INFORMATION:
NAME: Reiter, Stephen E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: 619-546-4737
TELEFAX: 619-546-9392
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CORRESPONDENCE ADDRESS:
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OPERATING SYSTEM:
SOFTWARE: Patentl
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CITY: Los Angeles
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444 South Flower Street, Suite 2000
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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15-JUN-1993
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US-09-851-896-84
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US-08-907-598-10/c
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US-08-907-598-10
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Best Local Similarity 78.3
Matches 18; Conservative
GENERAL INFORMATION:
APPLICANT: C. Frank Bennett
                                                                                                                                                                       Query Match
Best Local Similarity
Matches 18; Conser
                             Sequence 84, Application US/09851896 Patent No. 6410325
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MEDIUM TYPE: Floppy disk
COMPUTER: LEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 85:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 493-4935
                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION: NAME: Coruzzi, Laura A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                            MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                      TYPE:
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10036-2711
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Friedrich, Glenn
                                                                                                                                                                                                                                                                                                                                                                         (650) 493-5556
                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sands, Arthur
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Zambrowicz, Brian
                                                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PENNIE & EDMONDS LLP
                                                                                                                                                                                                                                                           DNA (genomic)
                                                                                                                                                                                                                                                                                        unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              800
                                                                                                                                                                                     57.78;
78.38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                57.7%;
78.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TARGETED GENE DISCOVERY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US/08/907,598
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                                                                                                                                                                         0;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                     8535-015-999
                                                                                                                                                                                      Score 15; DB 3;
Pred. No. 2.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 4;
                                                                                                                                                                                                     DB 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ر.
                                                                                                                                                                                                  Length 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
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                                                                                                                                                                         Gaps
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Susan M.

Freier

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                                                                                            ; MOLECULE TYPE: Other Nucleic Acid US-07-971-819A-32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-851-896-84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/851,896
CURRENT FILING DATE: 2001-05-08
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 84
                                Matches
                                               Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: ANTISENSE MODULATION OF PHOSPHOLIPASE A2, GROUP VI (CA2+-INDEPENDENTIALE OF INVENTION: EXPRESSION FILE REFERENCE: RTS-0220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Antisense Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                REGISTRATION NUMBER: 32,630
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (510) 814-2863
                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: WordPerfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 19930203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Gelfand, David H. APPLICANT: Lawyer, Frances C. APPLICANT: Stoffel, Susanne
                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                           Local
                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM:
SOFTWARE: WordPer
                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 07110-1199
                                                                                                                           TOPOLOGY:
                                                                                                                                       STRANDEDNESS:
                                                                                                                                                       TYPE: nucleic acid
                                                                                                                                                                                                                                     TELEFAX: (510)
                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Diskette, 3.50 inch, 800 Kb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Nutley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
                                                                                                                                                                       LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION:
                              l Similarity
19; Conserv
1 TGGTGCATGTCTTTGGTCCCAGCTAC 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CATTTCTTTAGTCCCAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CATGTCTTTGGTCCCAGC 23
                                                                                                                                                                                                                                                                                                                  Stacey R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New Jersey
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Andrew T. Watt
                                                                                                                                                                         36 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  340 Kingsland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                           linear
                                                                                                                                                                                                                                                                                                                                                                                                                                        Apple Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hoffmann-La Roche Inc.
                                                                                                                                     single
                                                                                                                                                                                                                                       814-2977
                                                                                                                                                                                                                                                                                      Sias, Fi... 32,630
                                                                                                                                                                                                                                                                                                                                                                                                                        Macintosh 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Polymerase Enzyme from Thermotoga Maritima
46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Purified Thermostable Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                56.9%;
88.9%;
                                                                                                                                                                                                                                                                                                                                                                          US/07/971,819A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Street
                                                                                                                                                                                                        32:
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Pred. No. 2.
                                           Score 14.8;
Pred. No. 2
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                                             .5e+02;
                                                            DB 1;
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                                                          Length 36;
                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                             0;
                           Gaps
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                             0;
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RESULT 7
US-07-977-434-28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Gelfan
APPLICANT: Abrams
                                                                                                                                                                                                                                                                                                                            FILING DATE: 22-AUG-1986
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 7.
FILING DATE: 15-AUG-1991
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 5
FILING DATE: 28-SEP-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: WordPerfect 2.1 CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy
                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 22-DEC-
                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 17-JUN-
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APPLICATION NUMBER:
FILING DATE: 15-MAY-
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APPLICATION NUMBER:
FILING DATE: 28-SEP-
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PRIOR APPLICATION DATA:
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                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 20-SEP-
                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
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                                                                                                                                                                        PRIOR APPLICATION DATA:
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ELECOMMUNICATION INFORMATION:
                                                                                                                                   APPLICATION NUMBER: US 609,157 FILING DATE: 02-NOV-1990
                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: WO PFILING DATE: 21-DEC-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: FILING DATE: 12-JAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: New Jersey
ZIP: 07110-1199
                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM:
              REGISTRATION NUMBER: 31 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28, Application US/07977434
5. 5466591
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TGAAGCATGTCTTTGTCACCGGTTAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             340 Kingsland Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Abramson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gelfand, David H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Macintos
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-SEP-1990
                                                                                                                                                                                                                                          JMBER: US 585,471
20-SEP-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-MAY-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           JMBER: US 590,213
28-SEP-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-JUN-1987
                                                                                                                                                                                          22-DEC-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-JAN-1988
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5' TO 3' EXONUCLEASE MUTATIONS
                                                                                                                                                                                                                                                                                                                                                                                                                   US 899,241
                                                                                                     US 557,517
                                                                                                                                                                                                          US 455,611
                                                                                                                                                                                                                                                                                                             WO PCT/US90/07641
                                                                                                                                                                                                                                                                                                                                                                US 746,121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US 063,509
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US 523,394
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US 590,466
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US 590,490
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US/07/977,434
              Case No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27
                5466591 8753
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NFORMATION FOR SEQ

(510) 814-2972 SEQ ID NO: 28

TELEPHONE:

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ANTI-SENSE:
US-07-977-434-28
                                    Query Match
Best Local Similarity
                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE: NO
                                                                                                                                                                          NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ent No.
                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 07/971,819
                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Diskette,
                                                                                         TOPOLOGY: li
                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS LENGTH: 36 nucleotide:
                                                                                                                                                             SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                   FILING'DATE: 03-FEB-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM: Macintosh SOFTWARE.
                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/0 FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                   TELEPHONE:
                                                                                                                                                                                                                                                      NAME: Stacey R. Sias, Ph.D. REGISTRATION NUMBER: 32,630
                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
                                                                                                                                              ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19;
                           19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            07110-1199
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                                                                                                                                  nucleic acid
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TGGTGCATGTCTTTGGTCCCAGCTAC 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/08475231
                                                                                                                                              36 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             340 Kingsland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gelfand, David H.
Lawyer, Frances C.
                          Conservative
                                                                                                                                                                                                     (510) 814-2977
                                                                                                                                                                                                                                                                                                                                                                                            WordPerfec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stoffel, Susanne
                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hoffmann-La Roche Inc
                                                                                                                                                                                                                   (510)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Jersey .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NO
                                                                                          Other Nucleic Acid
                                                                                                                                                                                                                                                                                                  03-FEB-1993
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                                                                                                                    single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  leotides
                                                                                                                                                                                                                   814-2863
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                56.9%;
73.1%;
                                     56.9%;
73.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Polymerase Enzyme from Thermotoga Maritima
                                                                                                                                                                                                                                                                                                                                                                                                      Macintosh 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Purified Thermostable Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                US/08/475,231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Street
                                                                                                                                                                          32:
                                                                                                                                                                                                                                                                                                                                                                                                                                3.50 inch, 800 Kb storage
                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                       Score 14.8; DB 1;
Pred. No. 2.5e+02;
0; Mismatches 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 14.8;
Pred. No. 2
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                                                   DB 1;
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                                                  Length 36;
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                       Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 28,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: WordPerfect 2.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                FILING DATE: 02-NOV-
PRIOR APPLICATION DATA:
                                                                                                FILING DATE: 22-DEC-:
PRIOR APPLICATION DATA:
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                                                                                                                                                    PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                         FILING DATE: 22-AUG-:
PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
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                                                                                                                                                                    APPLICATION NUMBER: US 5: FILING DATE: 20-SEP-1990
                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 7. FILING DATE: 15-AUG-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: New Jersey ZIP: 07110-1199
                                APPLICATION NUMBER:
                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 143,441 FILING DATE: 12-JAN-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 523,394 FILING DATE: 15-MAY-1990
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INFORMATION:
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22-AUG-1986
                                                                 02-NOV-1990
                                                                                                                   22-DEC-1989
                                                                                                                                                                                                                       21-DEC-1990
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28-SEP-1990
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                                                                                us 609,157
                                                                                                                                                                                                                                         WO PCT/US90/07641
                                US 557,517
                                                                                                                                   US 455,611
                                                                                                                                                                                                                                                                                           US 746,121
                                                                                                                                                                                                                                                                                                                                                                                               US 063,509
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                                                                                                                                                                                    US 585,471
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                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 523,394
                                                                                                                                                                                                                                                                                   FILING DATA:
PRIOR APPLICATION DATA:
OF THE PRIOR APPLICATION NIMBER: US 590,213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: Ca
TELECOMMUNICATION INFORMATION:
                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 746,121
FILING DATE: 15-AUG-1991
                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:

PRIOR APPLICATION NUMBER: US 590,490
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: WordPerfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
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HYPOTHETICAL:
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APPLICATION NUMBER:
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APPLICANT: Abramson, Richard
                                                                                                             PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 590,466 FILING DATE: 28-SEP-1990
                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 59 FILING DATE: 28-SEP-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 1400 Fifty-third Street CITY: Emeryville
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                                                                                FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: POFILING DATE: 19910930
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SOFTWARE: WordPerfect 5.0
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                                                                                               APPLICATION NUMBER:
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                                                                                                                            UMBER: US 063,509
17-JUN-1987
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73.1%;
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                                                                                             US 899,241
WO PCT/US90/07641
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Pred. No. 2.5e+02;
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Best Local
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           ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
                                                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                        SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                          APPLICATION NUMBER: US (FILING DATE: 21-APR-1989
                                                                                                                                                                                                                          OPERATING SYSTEM:
                                                                                                                                                                                                                                                        MEDIUM TYPE:
                                                                                                                                                           FILING DATE:
                                                                                                                                                                         APPLICATION NUMBER:
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US 07/341,562

US/08/222,177A

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US-08-222-177A-270
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                                                                                                                                                                                                                      Patent No. 5582979
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 5:
FILING DATE: 24-JUL-1990
                                            COMPUTER READABLE FORM:
                                                                                                                                                CORRESPONDENCE ADDRESS
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ANTI-SENSE:
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NAME: Sias Ph.D, Stacey R.
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                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                  NUMBER OF SEQUENCES:
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                                                                         COUNTRY:
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                                                                                                                                   ADDRESSEE:
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                                                                                                                                                                                                                                                   270, Application US/08222177A
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19; Conser
                                                                                                                                                                                                                                                                                                                                                       1 TGGTGCATGTCTTTGGTCCCAGCTAC 26
                                                           53717-1914
                                                                                                        Madison
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                                                                                                                                                                                                                                                                                                                             TGAAGCATGTCTTTGTCACCGGTTAC 27
                                                                                        Wisconsin
                                                                                                                    E: DeWitt Ross & Stevens, S.C. 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                           Weber,
                                                                         USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                               NO
                                                                                                                                                                                                                                                                                                                                                                                       Conservative
E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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02-NOV-1990
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                                                                                                                                                                                                           James L.
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73.1%;
                                                                                                                                                                          LENGTH POLYMORPHISMS IN (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS
                                                                                                                                                                 460
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                                                                                                                                                                                                                                                                                                                                                                                                   Score 14.8; DB 5;
Pred. No. 2.5e+02;
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RESULT 13
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US-08-239-256-18
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                                                                                                            Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: BOIME, IRVING
                                                                                                                                                                                                                                     NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO:
                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 29,
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 887-1500
                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION: NAME: MURASHIGE, KATE H.
                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: BOIME, IRVING
APPLICANT: MATZUK, MARTIN M.
APPLICANT: KEENE, JEFFREY L.
TITLE OF INVENTION: CTP EXTENDED FORM OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IMMEDIATE SOURCE:
CLONE: mfd78p2
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                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: FILING DATE: 06-MA CLASSIFICATION: 43
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                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE:
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18; Conserv
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                                                                 GTGCATGTCTTTGGTCCCAGCTAC 26
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Pred. No. 4e+02;
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Pred. No. 3.6e+02;
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; OTHER INFORMATION: Primer
US-09-286-959B-12
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US-09-286-959B-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: SEQ ID NO:20:
                                                                                                                                                                Sequence 9, Application US/08283857 Patent No. 5792742
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/080,783
PRIOR FILING DATE: 1998-04-06
NUMBER OF SEQ ID NOS: 24
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 12
                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
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Best Local S
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                                                                                                                                                   GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: TELOMERAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Johns Hopkins University
                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                          APPLICANT: BARON, MARTÍN
APPLICANT: CAMPBELL, Tain D.
APPLICANT: WILLIAMS, Michael, J.
TITLE OF INVENTION: FIBRONECTIN FIBRIN-BINDING PEPTIDES,
TITLE OF INVENTION: CODING THEREFOR AND USERS THEREOF
            NUMBER OF SEQUENCES: 1
                                                                                                                    APPLICANT: APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 01-JUN-1990 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA: APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: BOIME, IRVING; MATZUK, MARTIN M.
TITLE OF INVENTION: MODIFIED FORMS OF REPRODUCTIVE HORMONES
NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 41
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 ADDRESSEE:
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                                                                                                                    GOLD, Leslie I.
ROSTAGNO, Agueda A.
                                                                                                                                                                                                                                                                                                                   Conservative
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BROWDY AND NEIMARK
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75.0%;
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Pred. No. 4e+02;
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Pred. No. 4
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STREET:

419 Seventh Street, N.W.,

Suite 300

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CITY: Washington
STATE: D.C.

STATE: D.C.

STATE: D.C.

CONTRY: USA

ZIP: 20004

COMPUTER READABLE FORM:

MCOMPUTER READABLE FORM:

COMPUTER READA
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Result
No.
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Maximum Match
Listing first
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seq length: 50
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26
1 TGGTGCATGTCTTT
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/cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US09_NEW_PUB.seq:*
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/cgn2_6/ptodata/2/pubpna/US10_NEW_PUB.seq:*
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/cgn2_6/ptodata/2/pubpna/US60_NEW_PUB.seq:*
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/cgn2_6/ptcodata/2/pubpna/PCT_NEW_PUB.seq:*
/cgn2_6/ptcodata/2/pubpna/US06_NEW_PUB.seq:*
/cgn2_6/ptcodata/2/pubpna/US06_PUBCOMB.seq:*
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US-10-198-069-46
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US-09-860-738C-38
US-09-860-738C-38
US-09-860-738C-59
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Compugen
Sequence seq
                                 Sequence 10, Appl
Sequence 10, Appl
Sequence 10, Appl
Sequence 105, Appl
Sequence 127897,
Sequence 33, Appl
Sequence 38, Appl
Sequence 48, Appl
Sequence 50, Appl
Sequence 53, Appl
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Sequence 54, Appl
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US-09-964-667-10
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                                                                                 INFORMATION FOR SEQ ID NO: 10: SEQUENCE CHARACTERISTICS: LENGTH: 26 base pair:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Esmond, Robert W.
REGISTRATION NUMBER: 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS: ADDRESSEE: Sterne,
                                                                                                                                                                  REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 26 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                    TELEPHONE: 202-371-2600
TELEFAX: 202-371-2540
                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Washington
STATE: DC
                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Wands,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jack R.
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US-09-951-402-8

US-09-951-401-8

US-09-922-101-8

US-10-098-263B-1457

US-10-098-263B-105579

US-10-098-263B-121069

US-09-912-263-435

US-09-795-006A-4

US-09-780-164-1792
                                                                                                                                                                                                                                                              0 US-09-801-274-1391

US-10-001-876-109

US-10-091-876-30

US-10-098-2638-23537

US-110-098-2638-130235

US-10-098-2638-130235

US-09-801-274-1727

0 US-09-801-274-1727

0 US-09-877-156-11
  0 US-09-795-006A-13

US-10-198-069-39

US-10-098-263B-56376

US-10-098-263B-76416

US-10-098-263B-11162

US-10-098-263B-111673

US-10-098-263B-1127898

US-10-098-263B-127898
sequence 1727, Ap

Sequence 11, Appl

Sequence 11, Appl

Sequence 8, Appl

Sequence 8, Appl

Sequence 10579,

Sequence 10579,

Sequence 1721069,

Sequence 435, Appl

Sequence 43, Appl

Sequence 437, Appl

Sequence 437, Appl

Sequence 137, Appl

Sequence 1373, Sequence 116373,

Sequence 116373,

Sequence 117898,

Sequence 127898,
                                                                                                                                                                                                                                                                                                          Sequence 30, Appl
Sequence 23537, A
Sequence 130235,
                                                                                                                                                                                                                                                                                                                                      Sequence 109, Sequence 30, A
                                                                                                                                                                                                                                                                                                                                                                      Sequence 1391,
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ZIP: 20005-3934

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version
TOPOLOGY: linear MOLECULE TYPE: cDNA SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Transgenic Animals and Cel
Screening Drugs Effective
of Alzheimer's Disease
                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/09/964,667 FILING DATE: 28-Sep-2001 CLASSIFICATION; <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Sterne, Kessler, Goldstein STREET: 1100 New York Ave., Suite 600
                                                                                                                                                                                                                                                               32,893
                                                                                                                                                                                                                                        0609.4370000
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Treatment
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US-09-964-412-10
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                                                                                                                                                        RESULT 3
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US-09-964-666-10
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                                                                     Sequence 10, Application User Patent No. US20020129391A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 26
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/964,666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 0609.4370000 TELECOMMUNICATION INFORMATION:
                       TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 28-Sep-2001
CLASSIFICATION: <UNKnown>
ATTORNEY/AGENT INFORMATION:
NAME: Esmond, Robert W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Transgenic Animals and Cell Lines for Screening Drugs Effective for the Treatment or Prevention
                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                     1 TGGTGCATGTCTTTGGTCCCAGCTAC 26
                                                                                                                                                                                                                                                                                            26;
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                                                                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                               TGGTGCATGTCTTTGGTCCCAGCTAC 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Esmond, Robert W. REGISTRATION NUMBER: 32,893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TGGTGCATGTCTTTGGTCCCAGCTAC 26
                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 202-371-2540
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Sterne, Kessler, Goldstein & Fox, P.L.L.C
STREET: 1100 New York Ave., Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE:
                                                                                                                  Application US/09964412
                                                                                                                                                                                                                                                                                          Conservative
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                                       de la Monte, S
Wands, Jack R.
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Wands, Jack R.
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Transgenic Animals and Cell Lines for Screening Drugs Effective for the Treatment or Prevention
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                                                         Suzanne
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Pred. No. 0.0055;
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Pred. No. 0.0055;
                                                                                                                                                                                                                                                                                          Mismatches
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               ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: oligonucleotide US-10-198-069-46
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; MOLECULE TYPE: cDNA; SEQUENCE DESCRIPTION: SEQ ID NO: 10: US-09-964-412-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-10-198-069-46/c
                                                                                                                                                SOFTWARE: PatentIn Ver. SEQ ID NO 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 46, Application US/10198069 Publication No. US20030096756A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local
                                                                                                                                                                                      PRIOR FILING DATE: 2001-07-19
PRIOR APPLICATION NUMBER: 60/306,150
PRIOR FILING DATE: 2001-07-19
PRIOR APPLICATION NUMBER: 60/331,477
PRIOR FILING DATE: 2001-11-16
NUMBER OF SEQ ID NOS: 48
                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/10/198,069
CURRENT FILING DATE: 2002-07-19
                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: PEPTIDES EFFECTIVE IN THE TREATMENT OF TUMORS AND OTHER TITLE OF INVENTION: CONDITIONS REQUIRING THE REMOVAL OR DESTRUCTION OF TITLE OF INVENTION: CELLS
                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: 59003.000009
                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: AVERBACK,
                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/306,161
                                                                             FEATURE:
                                                                                         ORGANISM: Artificial Sequence
                                                                                                              TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: 202-371-26
TELEFAX: 202-371-2540
INFORMATION FOR SEQ ID NO: 10:
                                                                                                                                  LENGTH: 39
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26;
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TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Washington
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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Pred. No.
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ATTORNEY JAGEN 1 INFORMATION: <UNKNOWN>
ATTORNEY AGENT INFORMATION:

NAME: Esmond, Robert W.
REGISTRATION NUMBER: 32,893
REFERENCE JOCKET NUMBER: 0609.4370000
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/964,412
FILING DATE: 28-Sep-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox, P.L.L.C.
STREET: 1100 New York Ave., Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of Alzheimer's Disease
                       DB 10;
0.0055;
       0
                                               Length 26;
  Indels
  0;
Gaps
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                                          QΥ
                                                                                                                                                ; ORGANISM: Homo sapien US-10-098-263B-127897
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-893-238-105
                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 131066 .
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 127897
                                                                                Matches
                                                                                                 Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 105
LENGTH: 28
                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No. US20020150973A1
                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/098,263B
CURRENT FILING DATE: 2003-01-08
PRIOR APPLICATION NUMBER: 60/276,759
PRIOR FILING DATE: 2001-03-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Moore, K.
APPLICANT: Nagle, D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF BODY WEIGHT DISORDERS, INCLUDING
FILE REFERENCE: 7853-237
                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Mittman, Michael
TITLE OF INVENTION: Human Microarray
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 60/104,978
PRIOR FILING DATE: 1998-10-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/893,238
CURRENT FILING DATE: 2001-06-27
PRIOR APPLICATION NUMBER: 09/245,041
PRIOR FILING DATE: 1999-02-05
PRIOR APPLICATION NUMBER: 60/093,630
PRIOR FILING DATE: 1998-07-21
PRIOR FILING DATE: 1998-07-21
                                                                                                                                                                                   LENGTH: 25
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: 3118.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
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         105, Application US/09893238
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                                                                                Conservative
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78.3%;
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Pred. No. 8.
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Pred. No. 1.1e+03
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US-09-860-738C-48
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US-09-860-738C-35
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Publication No. US20030040620Al
GENERAL INFORMATION:
APPLICANT: Langmore, John
APPLICANT: Makarov, Vladimir
TITLE OF INVENTION: Method of Producing a DNA Library Using Positional Amplificati
FILE REFERENCE: UMIC:047US0/10103482
CURRENT APPLICATION NUMBER: US/09/860,738C
CURRENT FILING DATE: 2001-05-18
NUMBER OF SEQ ID NOS: 121
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 38
LENGTH: 23
TYPE: DNA
                                                                                                               GENERAL INFORMATION:
                                                                                                                              Sequence 48, Application US/09860738C Publication No. US20030040620A1
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APPLICANT: Langmore, John APPLICANT: Makarov, Vladimir APPLICANT: Makarov, Vladimir TITLE OF INVENTION: Method of Producing a DNA Library Using Positional Amplificati FILE REFERENCE: UMIC:047050/10103482 CURRENT APPLICATION NUMBER: US/09/860,738C CURRENT FILING DATE: 2001-05-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Langmore, John
APPLICANT: Makarov, Vladdimir
TITLE OF INVENTION: Method of Producing a DNA Library Using Positional Amplificati
FILE REFERENCE: UMIC:047US0/10103482
CURRENT APPLICATION NUMBER: US/09/860,738C
CURRENT FILING DATE: 2001-05-18
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TYPE: DNA
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FILE OF INVENTION: Method of Producing a DNA Library Using Positional Amplification FILE OF INVENTION: Method of Producing a DNA Library Using Positional Amplification CURRENT FILING DATE: 2001-05-18

NUMBER OF SEQ ID NOS: 121

SOFTWARE: Patentin version 3.1

SEQ ID NO 50
LENGTH: 23
TYPE: DNA
ORGANISM
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; Sequence 53, Application US/09860738C; Publication No. US20030040620A1
; GENERAL INFORMATION:
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; OTHER INFORMATION: Primer
U$-09-860-738C-50
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APPLICANT: Makarov, Vladimir
TITLE OF INVENTION: Method of Producing a DNA Library Using Positional Amplification
FILE REFERENCE: UMIC:047US0/10103482
CURRENT APPLICATION NUMBER: US/09/860,738C
CURRENT FILLING DATE: 2001-05-18
NUMBER OF SEQ ID NOS: 121
SOFTWARE: Patentin version 3.1
SEQ ID NO 53
LENGTH: 23
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Best Local Similarity
Matches 17; Conserv
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 48
LENGTH: 23
                                                                                                TYPE: DNA
ORGANISM: Unknown
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OTHER INFORMATION: Primer
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Pred. No. 1.3e+03;
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US-09-860-738C-55
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TITLE OF INVENTION: PEPTIDES EFFECTIVE IN THE TREATMENT OF TUMORS AND OTHER
                 APPLICANT: AVERBACK,
                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Unknown
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TYPE: DNA
ORGANISM: Unknown
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; OTHER INFORMATION: Primer US-09-860-738C-74
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Sequence 45, Application US/10198069 Publication No. US20030096756A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.1 SEQ ID NO 74
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APPLICANT: Makarov, Vladdmir
APPLICANT: Makarov, Vladdmir
TITLE OF INVENTION: Method of Producing a DNA Library Using Positional Amplificati
FILE REFERENCE: UNIC:047USO/10103482
CURRENT APPLICATION NUMBER: US/09/860,738C
CURRENT FILING DATE: 2001-05-18
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SOFTWARE: PatentIn version
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APPLICANT: Langmore, John
APPLICANT: Makarov, Vladimir
TITLE OF INVENTION: Method of Producing a DNA Library Using Positional Amplification
FILE REFERENCE: UMIC:047US0/10103482
CURRENT APPLICATION NUMBER: US/09/860,738C
CURRENT FILING DATE: 2001-05-18
NUMBER OF SEQ ID NOS: 121
SOFTWARE: Patentin version 3.1
SEQ ID NO 36!
LENGTH: 41
TYPE: DNA !
ORGANISM: Unknown
FORGANISM: Unknown
OTHER INFORMATION: Primer
US-09-860-738C-36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/306.161
PRIOR FILING DATE: 2001-07-19
PRIOR APPLICATION NUMBER: 60/306.150
PRIOR FILING DATE: 2001-07-19
PRIOR APPLICATION NUMBER: 60/331,477
PRIOR FILING DATE: 2001-11-16
NUMBER OF SEQ ID NOS: 48
SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 45
LENGTH: 39
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Search completed: June 22, 2003, 03:18:16
Job time : 78.7812 secs
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Best Local Similarity 81.0
Matches 17'; Conservative
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Best Local Similarity 81.0

Matches 17; Conservative
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CURRENT FILING DATE: 2002-07-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE: 59003.000009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         40 GGTGCATGTGTATCGTCCGAG 20
                                                                                            2 GGTGCATGTCTTTGGTCCCAG 22
                                                                                                                                                                  56:2%;
81.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   56.2%; Score 14.6; 81.0%; Pred. No. 1.4; tive 0; Mismatches
                                                                                                                                                 0;
                                                                                                                                                                Score 14.6;
Pred. No. 1.
                                                                                                                                               Mismatches
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L.4e+03;
                                                                                                                                                                  .4e+03;
                                                                                                                                                                                      DB 9;
                                                                                                                                                                                Length 41;
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                           Database :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Minimum DB seq length: 0 Maximum DB seq length: 50
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Perfect score:
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Gapop 10.0 , Gapext 1.0
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26
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gb_est1:*
gb_est2:*
gb_htc:*
gb_est3:*
gb_est4:*
em_gss_hum:*
em_gss_pin:*
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em_gss_yrt:*
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em_gss_fun:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

### SUMMARIES

C			O	C	ი	Result No.
σ	G	4	ω	N	1	
16.6	16.8	17	18	18	19.2	esult No. Score
63.8	64.6	65.4	69.2	69.2	73.8	Query Match
34	33	48	45	40	42	Query Match Length DB I
17	14	14	14	14	14	DB
AZ393855	н85333	N79917	T92229	N95034	H47110	DB ID
AZ393855 1M0157D07	H85333 ys90e04.rl	N79917 yz86a06.rl	T92229 ye17f11.r1	N95034 zb32b05.s1	H47110 yp77e03.s1	Description

# ALIGNMENTS

	COMMENT	TITLE		AUTHORS	REFERENCE		ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION		DEFINITION	LOCUS	RESULT 1 H47110/c
Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108 Tel: 314 286 1800 Fax: 314 286 1810 Email: est@watson.wustl.edu Insert Size: 704 High quality sequence starts: 1 High quality sequence stops: 1	Unpublished (1995) Contact: Wilson RK	.R., Williamson,A., Wohldmann,P. and Wilson,R. The WashU-Merck EST Project	,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston	Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman	1 (bases 1 to 42)	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.	Homo sapiens	human.	EST.	H47110.1 GI:923162	H47110	IMAGE:193468 3' similar to gb:Z13009_rna1 EPITHELIAL-CADHERIN PRECURSOR (HUMAN);, mRNA sequence.	oleen 1NFLS Homo sapi	H47110 42 bp mRNA linear EST 16-AUG-1995	

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REFERENCE
AUTHORS
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N95034/c
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SOURCE
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JOURNAL
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1 (bases 1 to 40)

Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P. and Wilson,R.

76. WashU-Merck EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                   N95034.1
EST.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              zb32b05.sl Soares_parathyroid_tumor_NbHPA Homo sapiens cDNA clone IMAGE:305265 3' similar to gb|m87890|HUMALCE124 Human carcinoma cell-derived alu RNA transcript, (rRNA); gb:M91159 !!!! ALU CLASS E WARNING ENTRY !!!! (HUMAN);, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21;
                                                                                               This close is available royalty-free through LLNL; contact the This Consortium (info@image.llnl.gov) for further information
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 TGGTGCATGTCTTTGGTCCCAGCTAC 26
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Insert Length: 704 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Source: IMAGE Consortium, LLNL
This clone is available royalt
                                                                                  Seq
                                                                                                                                                    Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                      Washington University School of Medicine
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Contact: Wilson RK
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Location/Qualifiers
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                                                                                    primer: ETPrimer
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Location/Qualifiers
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/organism="Homo sapiens"
/db_xref⇒"GDB:1248675"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone_lib="Soares fetal liver spleen lNFLS"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/db_xref="GDB:3762489"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone="IMAGE:193468"
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80.8%;
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Вþ Ş

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JOURNAL
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                                                                                                                                                                                                                                         Source: IMAGE Consortium, LLNL ;
This clone is available royalty-free through LLNL ;
IMAGE Consortium (info@image.llnl.gov) for further i
Trace considered overall poor quality
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Schellenberg, K., Soares, M.B., Tan, F., Thierry-Meg, J., Trevas Underwood, K., Wohldmann, P., Waterston, R., Wilson, R. and Marr Generation and analysis of £80,000 human expressed sequence Genome Res. 6 (9), 807-828 (1996)
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45 bp mRNA linear EST 22-MAR-199
yel7fll.rl Stratagene lung (#937210) Homo sapiens cDNA clone
yel7fll.rl Stratagene lung (#937210) Homo sapiens cDNA clone
IMAGE:118029 5' similar to gb:M81181 SODIUM/POTASSIUM-TRANSPORTING
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: est@watson.wustl.edu
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/organism="Homo sapiens"
/db_xref="GDB:486318"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="Organ: parathyroid gland; Vector: pT7T3D (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; 1st strand cDNA was primed with a Not I · oligo(dT)
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/dev_stage="adult"
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Pred. No. 1.1e+03;
0; Mismatches 5
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48 bp mRNA linear EST 02-APR-1996 yz86a06.rl Soares_multiple_sclerosis_2NbHMSP Homo sapiens cDNA clone IMAGE:289906 5' similar to gb|M87911|HUMALNE57 Human carcinoma cell-derived Alu RNA transcript, (rRNA); gb:X54150_rna1 IMMUNOCLOBULIN ALPHA FC RECEPTOR PRECURSOR (HUMAN); contains element
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: est@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (1995)
Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J. Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P. and Wilson,R.
The WashU-Merck EST Project
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Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman
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Location/Qualifiers
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double-stranded cDNA was size selected, ligated to Eco
                                                                                                                                                                                                                                                                                                                                                                                                                                      /db_xref="taxon:9606"
/clone="IMAGE:289906"
                                         /organism="Homo sapiens"
/db_xref="GDB:3905746"
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                                                                                                                                                                                                                                                                          /tissue_type="multiple sclerosis lesions"
/dev_stage="Age 46"
                                                                                                                                                                                                                                                                                                                                                                /sex="male"
                                                                                                                                                                                                                                                                                                                                                                                             /clone_lib="Soares_multiple_sclerosis_2NbHMSP"
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Pred. No. 1.2e+03
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                                                                                                                                                                                                                                                                                                                                                                                                                   High quality consortium, LLNL Source: IMAGE Consortium, LLNL; This clone is available royalty-free through LLNL; This clone is available royalty-free through LLNL; IMAGE Consortium (info@image.llnl.gov) for further i Trace considered overall poor quality tracert Length: 2596 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15 TGGTGTGCCTGTGATNCCAGCTAC 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi Mammalla; Eutherla; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 33)

Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holma,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterstran, Williamson,A., Wohldmann,P. and Wilson,R.

The WashU-Merck EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ys90e04.rl Soares retina N2b5HR Homo IMAGE:222078 5' similar to gb:Y00281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 TGGTGCATGTCTTTGGTCCCAGCTAC 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  High quality sequence starts: High quality sequence stops: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (1995)
Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    );, mk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: est@watson.wustl.edu
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                                                                                                                                                                                                                                                   /organism="Homo sapiens"
/db_xref="GDB:3850839"
/db_xref="taxon:9606"
/clone="IMAGE:222078"
/clone_lib="Soares retina N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                lesions from one patient was kindly provided by Dr. Kevin G. Becker (NINDS/NIH). "
a 7 c 18 g 15 t 1 others
                 double-stranded cDNA was size selected, ligated to Eco RI adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT7T3 vector
                                                                          the Not I and Eco RI sites of a modified pT7T3 vector (Pharmacia). Library went through one round of normalization to a Cot = 5. Library constructed by Bento Soares and M.Fatima Bonaldo. RNA from 4 multiple sclerosi
                                                                                                                                                                /tissue_type="retina"
/dev_stage="55 year old"
/lab_host="DH10B (ampicillin resistant)"
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                                                                                                                                                                                                                                    /sex="male"
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76.9%;
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The retinas were obtained from a 55
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gb:Y00281 RIBOPHORIN I PRECURSOR (HUMAN
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3.1e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     High
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0157 row: D column: 07
Seq primer: CGTTGTAAAACGACGGCCAGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Res
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34 bp DNA linear GSS 03-OCT-200
1M0157D07F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0157D07 F, DNA sequence.
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Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse whole genome scaffolding with paired end reads from 10kb
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adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gl14732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated
                                                                                                                   (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Caucasian and total cellular poly(A)+ RNA was extracted hrs after their removal. The retina RNA was kindly provided by Roderick R. McInnes M.D. Ph.D. from the University of Toronto. Library constructed by Bento Soares and M.Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                   /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Soares and M.Fatima Bonaldo. 9 c 9 g 7 t
                                                                                                                                                                                                                                                           Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                               /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGCIM0157D07"
                                                                                                                                                                                                                                                                                                   /Aab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                       /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                       /clone_lib="Mouse 10kb plasmid UUGC1M library"
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RESULT 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0332 row: K column: 15
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse whole مراجب , Mahmoud, M., Meenen, E., Pec, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Mouse whole مربية
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             l Similarity
19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert B. Weiss
University of Utah Genome
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dunn,D., Aoyagi,A., Bar
Islam,H., Longacre,S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1M0332K15F Mouse 10kb plasmid UUGC1M clone UUGC1M0332K15 F, DNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         High quality sequence stop: 48
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       308,
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10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gil4732141gblafr29072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated
                                                                                   was blunt end-repaired with T4 DNA polymerase and T4 polynucleotides kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                         (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA
                                                                                                                                                                                                                                          Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                               /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:10090"
/clone="UUGC1M0332K15"
                                                                                                                                                                                                                                                                                     /note="Vector:
                                                                                                                                                                                                                                                                                                                                /sex="Male"
                                                                                                                                                                                                                                                                                                                                                  /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                       /lab_host="E. Coli strain XL10-Gold, T1-resistant,
                                                                                                                                                                                                                                                                                                                                                                                                                      /strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Mus musculus
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with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to

The sheared, adaptored mouse DNA was

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ACCESSION
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AUTHORS
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T92842/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hillier, L., Lennon, G., Becker, M., Bonaldo, M.F., Chiapelli, B., Chissoe, S., Dietrich, N., DuBuque, T., Favello, A., Gish, W., Hawkins M., Hultman, M., Kucaba, T., Lacy, M., Le, M., Le, N., Mardis, E., Moore, B., Morris, M., Parsons, J., Prange, C., Rifkin, L., Rohlfing, T., Schellenberg, K., Soares, M.B., Tan, F., Thierry, Meg, J., Trevaskis, E., Underwood, K., Wohldmann, P., Waterston, R., Wilson, R. and Marra, M. Generation and analysis of 280,000 human expressed sequence tags Genome Res. 6 (9), 807-828 (1996)
                                                                                                                                                                                                                                                                                                                                                                                                             High quality sequence starts: 1 High quality sequence stops: 1 Source: IMAGE Consortium, LLNL This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Trace considered overall poor quality
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     T92842
Ye27a08.rl Stratagene lung (#937210) Homo IMAGE:118934 5' similar to gb:Z13009_rna1
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                                                                                                                                                                                                                                                                                                                                                                                                  Insert Length:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Washington University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97044478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mammalia;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Insert Size: 451
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314 286 1810
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Location/Qualifiers
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                                                     adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance:"

21 c 8 g 7 t
                                                                                                                                                                                                                                                  /db_xref="taxon:9606"
/clone="IMAGE:118934"
                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/db_xref="GDB:487223"
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                                                                                                                                                                                                                            /clone_lib="Stratagene lung (#937210)"
                                                                                                                                                                                                          'sex="male"
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82.6%;
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Pred. No. 4.6e+03;
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RESULT 9
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                                                                                  18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: est@watson.wustl.edu
Insert Size: 565
High quality sequence starts: 1 High quality sequence stops: 1.
Source: IMAGE Consortium, LLNL This clone is available royalty-free through LLNL: contact the IMAGE Consortium (info@image.linl.gov) for further information. Trace considered overall poor quality insert Length: 565 Std Error: 0.00
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Fax: 314 286 1810
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1 (bases 1 to 32)
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                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seq primer: M13RP1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRECURSOR (HUMAN);, mRNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CATGTCTTTGGTCCCAGCTAC
CATGCCTATGGTCCTAGCTAC
                                        CATGTCTTTGGTCCCAGCTAC 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                           9
                                                                                                                                                                                     /lab_host="SOLR cells (kanamycin resistant)"
/note="Organ: lung; Vector: pBluescript SR:, Site_1: EcoRI; Site_2: XhoI; Cloned unidirectionally. Primer: Oligo dT. normal lung. Average insert size: 1.0 kb; Uni-ZAP XR Vector; ~5' adaptor sequence: 5' GAATTCGGCACGAG 3' ~3' adaptor sequence: 5' CTCGAGTTTTTTTTTTTTTTTTT 3'"

a 10 c 8 g 5 t
                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Homo sapiens"
/db_xref="GDB:487271"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GI:724775
                                                                                                                                                                                                                                                                                                                                                                                                             /clone="IMAGE:118982"
                                                                                                                                                                                                                                                                                                                                                                     /sex="male"
                                                                                                                                                                                                                                                                                                                                                                                   /clone_lib="Stratagene lung (#937210)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                             'dev_stage="72 years"
                                                                                                      62.3%;
85.7%;
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                                                                                                      Score 16.2; DB 14
Pred. No. 6.2e+03;
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                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 St.
                                                                                                                          14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Louis, MO 63108
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sapiens cDNA clone
EPITHELIAL-CADHERIN
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RESULT 10

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BASE COUNT
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   RESULT 11
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LÓCUS
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                                                             22
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2M0110G21R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0110G21 R, DNA sequence.
                                                                                                                                         18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           High quality sequence stop: 24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Plate: 0110 row: G column: 21
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: 801 585 5606 Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  rukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 24)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AZ831249.1 GI:13001157 GSS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   and Wright, D., Weiss, R.
                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
ISLAM, ROSe,M., ROSe,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          house mouse.
                                                                                 TGGTGCATGTCTTTGGTCCCAG 22
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                                                                                                                                         Conservative
                                                                                                                                                                                                                                      9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ddunn@genetics.utah.edu
Length: 10000 Std Erro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
                                                                                                                                                                                                                                                                                                        was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gi14732114|gb|AF129072.1), a roovy number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
                                                                                                                                                                                                                                adaptored vector DNA, and transformed into chemically-competent E. coli XLLO-Gold (Stratagene) and selected for ampicillin resistance." a 6 c 5 g 4 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note-"Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Biomedical Polymers, Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:10090"
/clone="UUGC2M0110G21"
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/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /lab_host="E. Coli strain XL10-Gold, T1-resistant, F
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /sex="Male"
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                                                                                                                                                          60.0%;
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                                                                                                                                                       Score 15.6;
Pred. No. 1e
                                                                                                                                       Mismatches
                                                                                                                                                         1e+04;
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                                                                                                                                                                        DB 17;
                                                                                                                                       Indels
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DEFINITION
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                     KEYWORDS
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Best Local
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17; Conserv
human
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T94189 2
ye29h12.rl Stratagene lung
IMAGE:119207 5' similar to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: est@watson.wustl.edu
Insert Size: 320
High quality sequence starts: 1 High quality sequence stops: 1
Source: IMAGE Consortium, LLNL This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Trace considered overall poor quality
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   T90404

28 bp mRNA linear EST 20-MAR-:
yel6c02.rl Stratagene lung (#937210) Homo sapiens cDNA clone
lmaGE:117890 5', similar to gb:M73700 LACTOTRANSFERRIN PRECURSOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
(HUMAN);, mRNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Insert Length: 320
Seg primer: M13RP1
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                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:9606"
/clone="IMAGE:117890"
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/db_xref="GDB:486179"
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                                                                                                                                                                                                                                                                                                      58.5%;
85.0%;
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                                                                                                                                                                                                                                                                                                      Score 15.2;
Pred. No. 1
                                                                                                                                                                                                                                                                                Mismatches
                     28 bp mRNA linear EST 24-MAR-:

j (#937210) Homo sapiens cDNA clone

j (#937210) LACTOTRANSFERRIN PRECURSOR
                                                                                                                                                                                                                                                                                                        .6e+04;
                                                                                                                                                                                                                                                                                                                            DB
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RESULT
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                                                                          AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Insert Size: 348
Insert Size: 348
High quality sequence starts: 1 High quality sequence stops: 1
Source: IMAGE Consortium, LLML This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Trace considered overall poor quality Insert Length: 348

Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hillier, L., Lennon, G., Becker, M., Bonaldo, M.F., 'Chiapelli, B., Chissoe, S., Dietrich, N., DuBuque, T., Favello, A., Gish, W., Hawkins, M., Hultman, M., Kucaba, T., Lacy, M., Le, M., Le, N., Mardis, E., Moore, B., Mooris, M., Parsons, J., Prange, C., Rifkin, L., Rohlfing, T., Schellenberg, K., Soares, M.B., Tan, F., Therry Meg, J., Trevaskis, E., Underwood, K., Wohldmann, P., Waterston, R., Wilson, R. and Marra, M. Generation and analysis of 280,000 human expressed sequence tags Genome Res. 6 (9), 807-828 (1996)
              ,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., I
Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E.,
,R., Williamson,A., Wohldmann,P. and Wilson,R.
                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 33)

Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. 1
                                                                                                                                               Homo sapiens
                                                                                                                                                                                  EST
                                                                                                                                                                                                                                                   za50e12.s1 Soares fetal liver spleen 1NFLS Homo sapiens IMAGE:296014 3' similar to gb:X02067 H.sapiens mRNA for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tel: 314 286 1800 Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97044478
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Mammalia; Eutheria; Primates;
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                                                                                                                                                                                                   N67043.1
                                                                                                                                                                                                                               pseudogene (HUMAN);, mRNA sequence.
                                                                                                                                                                                                                                                                                        N67043
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/db_xref="GDB:487496"
/db_xref="taxon:9606"
/clone="IMAGE:119207"
                                                                                                                                                                                                   GI:1219168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /dev_stage="72 years"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     58.5%;
85.0%;
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Pred. No. 1.
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                                                     ., Marra, M., Parsons
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7SL RNA
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AZ438336.1
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                                                                                                                      Contact: Robert B. University of Utah University of Utah
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Contact: Wilson RK
                                                    Tel: 801 585 5606
Fax: 801 585 7177
                                                                                       84112,
                                                                                                                                                                            plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                         Mus musculus
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0228 row: E column: 1
                                                                                                                                                                                                                  Mouse whole genome
                                                                                                                                                                                                                               and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                               clone UUGC1M0228E19 F, DNA sequence.
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                                                                                                         308,
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                                                                                                        Biomedical
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/db_xref="GDB:1240935"
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/clone="IMAGE:296014"
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81.0%;
                                                                                                                                             Genome Center
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                Std Error:
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Di
Islam,H., Longacre,S., Mahmoud,M., Meenen,E.,
,M., Rose,M., Rose,R., Stokes,R., Tingey,A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Trace considered overall poor quality
Seq primer: ml3 -40 forward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Washington University School of Medicine 44444 Forest Park Parkway, Box 8501, St. 1
                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 38)
                                                                                                                                                                                                                                                                                                                                                             AZ438336 38 bp DN 1M0228E19F Mouse 10kb plasmid UUGC1M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  constructed by Bento Soares and M.Fatima Bonaldo." 9 c 11 g 7 t 1 others
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                                                                                                       scaffolding with paired
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 15.2; DB 14
Pred. No. 1.6e+04;
Research Bldg.,
                                                                                                                                                            Beacorn,T., Duval,B., Hamil,C.,
,M., Meenen,E., Pedersen,T., Reilly
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SLC,
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Best Local S
Matches 18
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A2489704 A2489704.1 GI:10659714
                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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High quality sequence stop: 38.
Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                           Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                        Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Welss,R.
                                                                                                                                               Rn.
                                                                                                                                                                                                                                    Unpublished (2000)
                                                                                                                                                                                                                                                               plasmid inserts
                                                                                                                                                                                                                                                                                Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                        308, Biomedical Polymers Research Bldg.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (9114732114 [95]AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli xL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                    USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Mus musculus
/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone="UUGC1M0228E19"
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78.3%;
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Pred. No. 2.1e+04;
0; Mismatches 5
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                                                                                                                                        2030 E.,
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Job time :

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Search completed: June 21,
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Class: plasmid ends
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                                                                                                                                                                                                                                                                                                                      adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gli4732114|gbhAF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli xL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0322L14"
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78.3%;
2003, 23:58:52
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Pred. No. 2.1e+04;
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SUMMARIES

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DEFINITION
Sequence 535 from Patent WO0206525.
ACCESSION
AX378746.1 GI:19574599
KEYWORDS
SOURCE
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
REFERENCE
AUTHORS
TITLE
JOURNAL
Patent: WO 0206525-A 535 24-JAN-2002;

PATENTAL

AND DNA
Linear PAT 18-MAR-2002

PAT 18-MAR-2002

PAT 18-MAR-2002

PAT 18-MAR-2002

AUTHORS
Cohen, D. Blumenfeld, M. Chumakov, I. Abderrahim, H. and Bihain, B.
Desity associated biallelic marker maps
JOURNAL
Patent: WO 0206525-A 535 24-JAN-2002;

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A84718
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Farinelli, i, and Mayer, P.
Farinelli, i, and Mayer, P.
METHOD OF NUCLEIC ACID SEQUENCING
Patent: WO 9844152-A 11 08-CCT-1998;
FARINELLI LAURENT (CH); MAYER PASCAL
                                                                                                                                                                                                                                                                                                                                                                                                          Ibd-related polymorphisms
Patent: WO 0142511-A 1707 14-JUN-2001;
WHITEHEAD INSTITUTE FOR BIOMEDICAL RES
Biotherapeutics Corporation (CA)
Location/Qualifiers
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Daly,M., Hudson,T.J., Lander,E.S., Rioux,J. and Siminovitch,K.
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11 from Patent WO9844152
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JP 2001017185-A/7.
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48 from Patent W00129257
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1 (bases 1 to 25)
Mizugaki,M. and Hiratsuka,M.
Mizugaki,M. and Hiratsuka,M.
Simple detection method of drug-metabolizing
Patent: JP 2001017185-A 7 23-JAN-2001;
OTSUKA PHARMACEUT CO LTD
OS Unidentified
ON JP 2001017185-A/7
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 47)
1 (bases 1 to 47)
Schork, N. and Skierczynski, B.
Methods of genetic cluster analysis and use thereof
Patent: WO 0129257-A 48 26-APR-2001;
GENSET (FR)
                                                                                       MICHINAO MIZUGAKI, MASAHIRO HIRATSUKA C12N15/09, C12Q1/68, C12Q1/68, C12N15/00
/organism='Unidentified'
Location/Qualifiers
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Sequence 37 from Patent EP1002861.
AX025434
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HOWELL STEVEN (GB); UNILEVER PLC (GB); LEDEBOER ADRIANUS MARINUS (NL); LOGT CORNELIS PAUL ERIK V D (NL); UNILEVER NV (NL); FRENKEN
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Sequence 84 from Patent W09923221.
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/note="LINKER"
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/db_xref="taxon:32644"
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/db_xref="taxon:32630"
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Synthetic eglin
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1 (bases 1 to 39)
Gill, P.T., Hussain, J.T. and Long, A.T.
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19; Conserv
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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1533 from Patent W00142511.
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/db_xref="taxon:32630"
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/db_xref="taxon:9606"
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Modified Eglin C sequence.
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Fritz,H., Maerki,W. and Alkan,S.
Process for the preparation of protease inhibitors
Patent: EP 0146785-A 12 03-JUL-1985;
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A34738
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  Fritz,H.,
   Rink, H., Liersch, M., Fritz, H., Maerki, W.
                                                                                      A34747.1
                                       unclassified
                                                                                                                                                                                                                                                                                                                                                      CIBA-GEIGY AG; PLANTORGAN WERK Heinrich G.E. Christensen
                                                                                                                                                                                                                                                                                                                                                                                                                                            unidentified
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1 (bases 1 to 41)
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9 c 7 g
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                                                                                      GI:21694241
                                                                                                                                                                                                                                                                                                                    /organism="unidentified"
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79.2%;
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72.4%;
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             Sieber, P.,
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Pred. No.
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               Meyer, F.,
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               Seemueller, U.
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Cargill,M., Ireland,J.S. and Lander,E.S.
                                                                     Method and reagent for the inhibition of erg
Patent: WO 0188124-A 3417 22-NOV-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); GLAXO
                                                                                                                                                          synthetic construct
synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human single nucleotide polymorphisms
Patent: WO 0166800-A 1110 13-SEP-2001;
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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/note="Enzymatic Nucleic Acid"
8 c 10 g 12 t
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/db_xref="taxon:9606"
11 c 4 g
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/db_xref="taxon:32644"
7 c 9 g 1
                                                                                                                      von Carlowitz, I., Mcswiggen, J.A.,
                                                                                                                                              sequences.
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77.38;
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ORIGIN

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RESULT 15
AX114375
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DEFINITION
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VERSION
KEYWORDS
SOURCE
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AUTHORS
TITLE
JOURNAL
Search completed: June 21, 2003, 22:14:32 Job time: 606.344 secs
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Best Local Similarity
Matches 20; Conserv
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                                                                                                                                                                                                                          variation
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Sequence 44 from Patent W00129257.
AXII4375
AXII4375.1 GI:14031339
                                                                                                                                                                                                                                                                                                    Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 47)
Schork,N. and Skierczynski,B.
Methods of genetic cluster analysis and use thereof
Patent: WO 0129257-A 44 26-APR-2001;
                                                                                                                                                                                                                                                                                                GENSET (FR)
                                                                                                                  50.7%; Score 15.2; DB 6; ilarity 66.7%; Pred. No. 5.3e+04; Conservative 1; Mismatches 9;
                                                                                                                                                                                       /note="99-1572-440 : polymorphic base C or T" ll a 14 c 14 g 7 t 1 others
                                                                                                                                                                                                                        /organism="Homo sapiens"
/db_xref="taxon:9606"
24
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1. .47
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GenCore version 5.1.6 Copyright (c) 1993 - 2003 Compugen Ltd

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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    June 21, 2003, 21:23:51; Search time 164.531 Seconds
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/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1981
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/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA2002
                                                                   /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA2000.DAT:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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	15.6	5	5	16	16	16	٠		16.4			16.6			$\vdash$	17.2				17.4	17.4	17.4	17.4							18	œ	œ	8	•	18.6
. 2	52.0	2	2	ω	ω.	ω.	4.	4.	4.	4.	5	5	5	6.	6	7.	7.	8	8	8	8	8	8	8	8	8	8	9.	9.	0	0	0	0		2
34	33	33	31	41	41	33	3 3	22	41	39	40	29	24	25	29	ω ω	29	50	47	47	41	41	41	41	33	41	41	33	. 24	29	34	ω ω	<u>ω</u>	29	47
22	24	24	22	24	24	24	24	21	24	22	22	21	24	22	21	24	21	22	22	21	24	24	24	24	22	24	24	24	22	21	22	24	24	21	24
AAH46543	w	ABK14188	AAI30622	ω	193	132	ABL49773	AAD00662	ABQ77547	AAF74239	AAH91207	AAD00661	ABA02134	AAF84351	AAA04503	ABN84112	AAA04311	AAL28723	AAF89248	AAZ66299	ABA96813	ABA96812	AAL43827	AAL43826	AAI65108	ABA94081	ABA94080	ABA01398	AAH91382	131	AAH27605	ABK14174	ABK12114	AAA04485	ABK41287
																																	٠		
DNA polymerase 17	Human pax protein	Human splicing fac	n single nu		cotrophin re		n tyr	hetic oligonu	3	analy	Human inflammatory	Synthetic oligónuc	Human zinc ion tra	æ	Polymorphic fragme	Human protein 11.1	Polymorphic fragme	_	e member cl			uteroglobi	oncogene	_	hypoxia-ind	multi-coppe	multi-	NF-	3	orph	II amir	in like gr	. 57	orphic fragm	Human USF2 gene bi

## ALIGNMENTS

RESULT 1 AAA03983

AAA03983 standard; DNA;

29 ВP

Polymorphism; hypertension; agammaglobulinemia; diabetes insipidus; Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome; Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis; 07-MAY-1998; 03-MAY-1999; progressic kidney disease; von Willebrands disease; forensic; tuberous scherosis; hereditary hemorrhagica telangiectasia; familial colonic polyposis; osteogenesis imperfecta; porphyria; Ehlers-Danlos syndrome; ss. (AFFY-) AFFYMETRIX INC. 07-MAY-1999; EP955382-A2 AAA03983; Homo sapiens. Polymorphic fragment of hypertension associated gene APOC3. 22-MAY-2000 (first entry) 10-NOV-1999 98US-0084641. 99US-0304232. 99EP-0250150 forensic; human;

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ABA01400/c
ID ABA014
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention provides polymorphic fragments of genes associated with hypertension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension with hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to find the precise role of a polymorphism in the disease, and this can be used to identify potential drugs which combat the disease. The polymorphisms can be tested for association with other diseases e.g. agammaglobulinemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, Wiskott-Aldrich syndrome, Fabrys disease, familial hypercholesterolemia, polycystic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     kidney disease, hereditary spherocytosis, von Willebrands disease, tuberous sclerosis, hereditary hemorrhagica telangiectasia, familial colonic polyposis, Ehlers-Danlos syndrome, osteogenesis imperfecta, a acute intermittent porphyria. The polymorphic forms can also be used forensics to identify individuals.
  Human NF-E2-associated factor in diagnosis and treatment of
                                                                     WPI; 2002-025846/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; NF-E2-associated factor 13; cytostatic; virucidal; immunomodulatory; antiinflammatory; haemostatic; gene therapy; haemopathy; HV infection; immunological disease; inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABA01400;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABA01400 standard; DNA;
                                                                                                                                                                                                                                                          19-MAR-2001; 2001WO-CN00368
                                                                                                                                                                                                                                                                                                         11-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human NF-E2-associated factor 13 probe #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-JAN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                 (BIOW-)
                                                                                                                                                                                                            22-MAR-2000; 2000CN-0115024
                                                                                                                                                                                                                                                                                                                                                  WO200175024-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                          developmental disorder; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel nucleic acids
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22; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 9 A; 11 C; 5 G; 3 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22; 53pp;
                                                                                                                                                                 GENE
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88.0%;
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                                                                                                                                                                 DEV
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Pred. No. 1
13 and
e.g. ma
                                                                                                                                                               SHANGHAI
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nd encoded
malignant
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polynucleotide, applicable tumor, hemopathy, HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21;
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1 ATCAACCTGGCGAACATGGTGAACCCC

Conservative

Indels

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Gaps

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RESULT 3
aBa01401/c
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AC ABa014
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AC ABa014
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DT 31-JAN
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Best Local S
Matches 23
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Best Local :
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                                                                                                                          The present invention relates to human Nr LL LUCY LAMS 2544). The factor and its coding sequence are useful in the AAMS 2544). The factor and its coding sequence are useful in the AAMS 2544). The factor and its coding sequence are useful in the AAMS 2544). The factor and developmenta
                                                                                                     and treatment of malignant tumours, haemopathy, HI immunological diseases, various inflammations and disorders. The present sequence is a probe, which
                                                                                                                                                                                                                                   Human NF-E2-associated factor 13 and encoded polynucleotide, applicable in diagnosis and treatment of e.g. malignant tumor, hemopathy, HIV infection, immunological diseases and inflammation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAM52544). The factor and its coding sequence are useful in the and treatment of malignant tumours, haemopathy, HIV infection, immunological diseases, various inflammations and developmenta disorders. The present sequence is a probe, which was used in
                                                                                                                                                                                                    Example 6; Page 15; 34pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABA01401 standard;
                                                              Sequence 41 BP; 7
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                                                                                                                                                                                                                                                                                                                                                                                            22-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                         19-MAR-2001; 2001WO-CN00368
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200175024-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                developmental disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; NF-E2-associated factor 13; cytostatic; virucidal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31-JAN-2002
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                                                                                                                                                                                                                                                                                                                                                             (BIOW-) BIOWINDOW GENE DEV INC SHANGHAI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             haemopathy; HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            infection,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lmmunomodulatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NF-E2-associated factor 13 probe #2
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                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                            2000CN-0115024
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             infection; immunological disease; inflammation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15; 34pp; Chinese
                                                                                             present invention
                                                              A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; 11 C; 13 G; 10 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         entry)
                68.7%;
85.2%;
                                                              11 C; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           68.7%;
85.2%;
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; Score 20.6; D; Pred. No. 30; 0; Mismatches
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Pred.
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                                                                                                                                                                       to human NF-E2-associated factor 13
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                                                              10 T; 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inflammation
                                                              other;
                                DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gene therapy; tumour;
                              Length
                                                                                                                        developmental
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                                 41;
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                                                                                                                                                       the
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                                                                                                                                                       diagnosis
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Дb

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RESULT 5
ABL49775/c
ID ABL49775 standard; D
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AC ABL49775;
XX
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AAS15589
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                                                                                                                                                                                                                                      В
                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                   The invention relates to an isolated polypeptide of human DNA mismatch repair protein 10. The polypeptide can be used for screening mimics, agonists, antagonists or inhibitors, or in peptide fingerprinting identification. The polypucleotide can be used as primers for nucleic acid amplification reactions, as probes for hybridisation reactions, or in producing gene chips or microarrays. Drug compositions, which contain the polypeptide, polypucleotide, mimics, agonists, antagonists, inhibitors and their preparations, can be used treatment and diagnosis of diseases relating to the polypeptide. In particular, the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mao
                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                mismatch repair protein 10.
                                                                                                                                                                                                                                                                             polypeptide and encoded polynucleotide are applicable in diagnosis and treatment of malignancy, haemopathy, human immunodeficiency virus (HTV) infection, immunological diseases and various inflammations. The present sequence represents PCR primer #2 used to amplify human DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 4; Page 13; 36pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New human DNA mismatch repair protein 10 for diagnosing and treating malignancy, haemopathy, human immunodeficiency virus infection, immunological diseases and inflammation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-MAR-2000; 2000CN-0115057
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-MAR-2001; 2001WO-CN00337
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; DNA mismatch repair protein 10; cytostatic; virucidal;
immunomodulatory; antiinflammatory; haemostatic; anti-HIV; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human DNA mismatch repair protein 10, PCR primer #2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BIOW-) BIOWINDOW GENE DEV INC SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunological disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human
                                                                                                                                                                                        Local
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                                                                                                                    TCAGCCTGGCCAACATGGTGAAAC
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                                                                                                                                                                                                                                     BP; 9 A;
                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                        64.0%;
                                                                                                                                                                                                                                     10 C; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     virus; malignancy; haemopathy; infection;
PCR primer; ss.
                                              41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     \omega
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВP
                                                                                                                                                                            0;
                                                                                                                                                                          Score 19.2; DB 24;
Pred. No. 1.1e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                     G;
                                                                                                                                                                                                                                     6 T; 0 other;
                                                                                                                   \omega
                                                                                                                                                25
                                                                                                                                                                                                        DB 24;
                                                                                                                                                                                                        Length
                                                                                                                                                                            Indels
                                                                                                                                                                                                        33;
                                                                                                                                                                          0;
                                                                                                                                                                          Gaps
                                                                                                                                                                          0;
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Ω
PR XX
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ABL49776/c
ID ABL49776 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       В
                                                                                                                   ΧX
                                                                                                                                                                      PN
                                                                                                                                                                                                        SS
                                                                                                                                                                                                                                           \Sigma \Sigma
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            invention also described a method for preparing (I) using DNA recombination techniques. (I) and the polynucleotide encoding it can be used in the treatment of diseases such as cancer and human immunodeficiency virus (HIV) infection. The present sequence represents a probe for human tyrosinase 10.34, which is used in an example from
                                                                                                                                                                    CN1325972-A.
                                                                                                                                                                                                                                         Human; tyrosinase;
HIV infection; can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 41 BP; 5 A; 12 C; 13 G; 11 T;
                                                                 31-MAY-2000; 2000CN-0116261
                                                                                                   31-MAY-2000;
                                                                                                                                     12-DEC-2001
                                                                                                                                                                                                                                                                                           Human tyrosinase 10.34 probe 2 SEQ ID NO:9
                                                                                                                                                                                                                                                                                                                                29-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes human tyrosinase 10.34 (I).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 6; Page 19 (Disclosure); 32pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New polypeptide-tyrosinase 10.34\ \text{for treating diseases} such as cancer and human immunodeficiency virus infection -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-196693/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             мао у,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31-MAY-2000; 2000CN-0116261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-MAY-2000; 2000CN-0116261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human
                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HIV infection; cancer; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-MAY-2002
                              (BODE-) BODE GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (BODE-) BODE GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tyrosinase 10.34 probe 1 SEQ ID NO:8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 ATCAACCTGGCGAACATGGTGAACCCC 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Xie Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ACCAGCCCGGCCAACATGGTGAAACCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                   2000CN-0116261
                                                                                                                                                                                                                                         cancer; probe; ss
                              DEV CO LTD SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DEV CO LTD SHANGHAI
                                                                                                                                                                                                                                                         enzyme; human immunodeficiency virus infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          enzyme; human immunodeficiency virus infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         63.3%;
                                                                                                                                                                                                                                                                                                                                                                                                  41
                                                                                                                                                                                                                                                                                                                                                                                                  ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19;
No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           .4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          <u>ر</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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RESULT 7
AAZ69526/c
          γQ
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Best Local S
Matches 22
     AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the
                                                               Claim
                                                                                                                    WPI; 2000-013267/01
                                                                                                                                          Cohen
                                                                                                                                                                                                                                                                WO9954500-A2
                                                                                                                                                                                                                                                                                                                      Key
                                                                                                                                                                                                                                                                                                                                                                diagnosis; single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                     Human map-related biallelic marker SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ69526 standard;
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                                                                                                                                                              (GEST ) GENSET
                                                                                                                                                                                    23-NOV-1998;
                                                                                                                                                                                               21-APR-1998;
                                                                                                                                                                                                                    21-APR-1999;
                                                                                                                                                                                                                                          28-OCT-1999
                                                                                                                                                                                                                                                                                                                                            Homo
                                                                                                                                                                                                                                                                                                                                                                                   Human genome;
genomic map;
                                                                                                                                                                                                                                                                                                                                                                                                                                            10-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ69526;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes human tyrosinase 10.34 (I). The present invention also described a method for preparing (I) using DNA recombination techniques. (I) and the polynucleotide encoding it can be used in the treatment of diseases such as cancer and human immunodeficiency virus (HIV) infection. The present sequence represents a probe for human tyrosinase 10.34, which is used in an example from
                                                                                                                                                                                                                                                                                                            variation
                                                                                                                                                                                                                                                                                                                                                                          haplotyping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 6; Page 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New
                                                                                el biallelic markers
of the human genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       present
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                                                                                                                                                                                                                                                                                                                                          sapiens
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for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   41
                                                           Page 1058; 2745pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACCAGCCCGGCCAACATGGTGAAACCC 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATCAACCTGGCGAACATGGTGAACCCC 27
                                                                                                                                        Blumenfeld
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                immunodeficiency virus infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention.
                                                                                                                                                                                                                                                                                                                                                                        biallelic marker; high density disequilibrium map;
haplotype; phenotype; polymorphic base; genotyping;
hybridiaation; identification; characterisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                           (first
                                                                                                                                                                                   98US-0082614
98US-0109732
                                                                                                                                                                                                                    99WO-IB00822
                                                                                                                                                                                                                                                                                              Location/Qualifiers replace(24,T)
/*tag= a
                                                                                                                                                                                                                                                                                   /standard_name=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (Disclosure); 32pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                           entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               63.3%;
81.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <u>ე</u>
                                                                                             used to
                                                                                                                                        Chumakov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ű
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10.34 for treating diseases
virus infection -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВP
                                                             English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 19;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <u>ი</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                             construct a
                                                                                                                                                                                                                                                                                    "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Т;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                other;
                                                                                                                                                                                                                                                                                                                                                                                                                      NO:3882
                                                                                                                                                                                                                                                                                                                                                                SNP;
                                                                                             high
                                                                                                                                                                                                                                                                                                                                                                ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 41;
                                                                                             density disequilibrium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            such
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cancer
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RESULT 8
AAK65537/c
ID AAK655
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20202020202020
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Best Local S
Matches 22
                   11-JUL-2000
14-JUL-2000
26-JUL-2000
26-JUL-2000
14-AUG-2000
14-AUG-2000
14-AUG-2000
14-AUG-2000
14-AUG-2000
14-AUG-2000
14-AUG-2000
14-AUG-2000
14-AUG-2000
                                                                                                                                                         30-JUN-2000;
07-JUL-2000;
07-JUL-2000;
11-JUL-2000;
                                                                                                                                                                                                                                                     02-MAR-2000;
16-MAR-2000;
17-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment.
                                                                                                                                                                                                        07-JUN-2000;
28-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                           Human; immune; ha cytostatic; gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N.B. The SEQ ID NOS 2852, 2913, and 3367, are not actually giver
                                                                                                                                                                                                                                                                                       24-FEB-2000;
                                                                                                                                                                                                                                                                                                                                    17-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-NOV-2001- (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAK65537 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 47 BP;
                                                                                                                                                                                                                                19-MAY-2000
                                                                                                                                                                                                                                           18-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAK65537;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              immune/haematopoietic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 ATCAACCTGGCGAACATGGTGAACCCC 27
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        2000US-0217496.
2000US-0218290.
2000US-0220964.
2000US-0224519.
2000US-0224519.
2000US-0225214.
2000US-0225214.
2000US-0225267.
2000US-0225268.
2000US-0225268.
2000US-0225268.
                                                                                                                                                                                2000US-0186350.
2000US-0189874.
2000US-0199076.
2000US-0199123.
2000US-0205515.
2000US-020467.
2000US-0214846.
2000US-0215135.
2000US-0216647.
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                                                                                                                                                                                                                                                                                                                                     2001WO-US01354
                                                                                                                                                          2000US-0216880
2000US-0217487
                                                                                                                                                                                                                                                                                       2000US-0184664
                                                                                                                                                                                                                                                                                                                                                                                                                                         haematopoietic;
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                                                                                                                                                                                                                                                                                                                                                                                                                           therapy; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; 14 C; 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   actually given a sequence in the Sequence Listing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA; 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 63.3%;
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   G; 13 T;
                                                                                                                                                                                                                                                                                                                                                                                                                              immune/haematopoietic antigen; cancer;
ine; metastasis; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              genomic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequence
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well as other
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14 - SEP - 2000;
21 - SEP - 2000;
21 - SEP - 2000;
21 - SEP - 2000;
25 - SEP - 2000;
26 - SEP - 2000;
27 - SEP - 2000;
29 - SEP - 2000;
29 - SEP - 2000;
20 - CCT - 2000;
02 - CCT - 2000;
02 - CCT - 2000;
03 - CCT - 2000;
04 - CCT - 2000;
05 - CCT - 2000;
07 - CCT - 2000;
08 - NOV - 2000;
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11-AUG-2000
18-AUG-2000
22-AUG-2000
22-AUG-2000
23-AUG-2000
23-AUG-2000
01-SEP-2000
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14-SEP-2000;
14-SEP-2000;
14-SEP-2000;
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14-SEP-2000;
                                           2000US-0246476.
2000US-0246477.
2000US-0246478.
2000US-0246478.
2000US-0246523.
2000US-0246524.
2000US-0246525.
2000US-0246525.
2000US-0246525.
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2000US-0241808.
2000US-0241809.
2000US-0241826.
2000US-0244617.
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2000US-0232401.
2000US-0233063.
2000US-0233064.
2000US-0233065.
2000US-0234224.
2000US-0234274.
2000US-0234274.
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2000US-0230438.

2000US-0231242.

2000US-0231244.

2000US-0231244.

2000US-0231244.

2000US-0231413.
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2000US-0239935.
2000US-0239937.
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2000US-0251868
2000US-0251869
                                                                                                                                                                                                                                                              SCI INC
                                                                                                                                                                                                                                             Ruben
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Nucleic acids encoding human immune/hematopoietic antigen polypeptides, useful for preventing, diagnosing and/or treating cancers and metastasis -

Disclosure; SEQ H Ö 20349; 3071pp + Sequence Listing; English

AAK54951 to AAK64702 encode the human immune/haematopoietic antigen (I) amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic activity, and can be used in gene therapy and vaccine production. (I), proteins and polynucleotides may be used in the prevention, diagnosis an treatment of diseases associated with inappropriate (I) expression. For example, they may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) polynucleotides may be used to produce the secreted (I), by inserting the nucleic acids into a host cell and culturing the cell to express the protein. (I) proteins and polynucleotides may be used to prevent, diagnose and treat immune/haematopoletic-related diseases, especially cancers and cancer metastases of haematopoletic-derived cells. AAK64703 to AAK87694 represent human immune/haematopoletic antigen genomic sequences from the present invention. AAK84942 to AAK5990 and AAM82169 represent sequences used in the exemplification of the present invention. present invention and

Sequence 49 BP; 11 A; 12 C; 13 G; 13 T; 0 other;

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Matches
        Query Match
Best Local 9
22; Conservative
       63.3%;
0;
        Score 19; DB 22;
Pred. No. 1.4e+02;
Mismatches
               Length 49,
Indels
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RESULT 9
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114-AUG-2000;
01-SEP-2000;
06-SEP-2000;
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13-OCT-2000;
08-NOV-2000;
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05-DEC-2000;
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                       Novel plas
treating,
proteins,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; nootropic; neuroprotective; cytostatic; dermatological; virucide; immunosuppressive; antiinflammatory; anti-HIV; antibacterial; vulnerary; antiparkinsonian; antisickling; antianaemic; antiarthritic; cancer; antiiparkinsonian; antisickling; artianaemic; antiinflammatory; antiinheumatic; hepatotropic; cerebroprotective; antiinflammatory; antiallergic; antidiabetic; antiulcer; anticonvulsant; antifungal; antiparasitic; cardiant; immune disorder; cardiovascular disorder; antionyological disease; infection; nephrotropic; gene therapy; vaccine; neurological disease; infection; nephrotropic; gene therapy; vaccine;
  Example
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04-FEB-2000;
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                       plasma membrane associated proteins useful
ing, preventing and/or prognosing disorders
ins, including cancer, immune response and o
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NO 233; 532pp + Sequence Listing; English.
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neuronal disorders
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The invention relates to a set of novel map-related biallelic markers, preferably located on obesity disorder-associated chromosomal regions chromosomes 3, 10 and 19. The markers are useful for genotyping or estimating the frequency of an allele in a population, for detecting a
                                                                                                                                                                                                                                                                                                                                     Set of novel map-related biallelic markers, preferably located on obesity disorder-associated chromosomal regions on chromosomes 3, obesity disorder. associated chromosomal regions on chromosomes 3, obesity disorder. The detecting statistical correlations between the correlations of the correlations of the correlations between the correlations of the correlations 
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                                                                                                                                                                                                                                                                                                          marker allele and a phenotype
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                  The invention provides polymorphic fragments of genes associated with hypertension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension diagnostic assays. Where the polymorphisms have strong correlation with hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to find the precise role of a polymorphism in the disease, and this can be used to identify potential
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            insulin disorders, atheromatous disease and cardiac insufficiency. The markers are useful for detecting a statistical correlation between a biallelic marker allele and a phenotype and/or between a biallelic mark haplotype and a phenotype. Sequences ABK41284-ABK41287 represent human LSR gene or USF2 gene biallelic markers.
                                                                                                                                                                                                                                                                                                         Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (AFFY-)
(UYCA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-MAY-1998;
03-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Polymorphism; hypertension; agammaglobulinemia; diabetes insipidus; Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome; Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis; polycystic kidney disease; von Willebrands disease; forensic; human; tuberous sclerosis; hereditary hemorrhagica telangiectasia; familial colonic polyposis; osteogenesis imperfecta; porphyria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA04485 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        association between a genotype or haplotype and a phenotype, e.g. a disease involving drug responses, obesity or disorders related to obesity, such as hyperuricaemia, digestive pathology, hepatic function disorders, cancer, cardiovascular disease, hypertension, hyperlipidaem
                                                                                                                                                                                                                                                                                                                                                     hypertension
                                                                                                                                                                                                                                                                                                                                                                             Novel nucleic acids containing polymorphisms used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EP955382-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Polymorphic fragment of hypertension associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ehlers-Danlos
                                                                                                                                                                                                                                                                                                                                                                                                                                  2000-107928/10
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UNIV CASE WESTERN RESERVE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                      Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chakravarti A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ACCATCMTGGCTAACACGGTGAAACCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first
                                                                                                                                                                                                                                                                                                      38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98US-0084641.
99US-0304232.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99EP-0250150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              62
77
  disease. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         . 0.9
                                                                                                                                                                                                                                                                                                   English.
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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No. 2.
polymorphisms can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene PGIS
                                                                                                                                                                                                                                                                                                                                                                             in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                          the
be
                                                                                                                                                                                                                                                                                                                                                                          diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hyperlipidaemia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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Ωy

ATCAACCTGGCGAACATGGTGAA

Query Match Best Local Matches

Similarity

60.7%; 87.0%;

Score 18.2;

Conservative

0;

Mismatches

Pred.

NO.

9e+02; DB

24;

Length Indels

33;

0;

Gaps

0;

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RESULT 12
ABK12114/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                        The invention relates to an isolated polypeptide of human EHD 9.57, the cDNA encoding it, and its fragment, analogue or derivative. Also included are vectors expressing the protein, a host cell comprising the vector, the isolation of modulators of the protein and an anti-EHD antibody. The protein and nucleic acid are used in diagnosis and treatment of a fetation abnormality, tumour, immune deficiency, neural system dysfunction disease, and protein metabolism disorders. The present sequence is a PCR primer used to clone the cDNA encoding the EHD protein (not defined).
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; ss; PCR; EHI immune deficiency;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 colonic polyposis, Ehlers-Danlos syndrome, osteogenesis imperfecta, and acute intermittent porphyria. The polymorphic forms can also be used in forensics to identify individuals.
 Sequence 33
                                                                                                                                                                                          Human EHD protein 9.57 and treatment of -
                                                                                                                                                                                                                                   WPI; 2002-172121/22
                                                                                                                                                                                                                                                              мао у,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         kidney disease, hereditary spherocytosis, von Willebrands disease, tuberous sclerosis, hereditary hemorrhagica telangiectasia, familial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       syndrome, Fabrys
kidney disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  association with other diseases e.g. agammaglobulinemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, Wiskott-Aldrich syndrome, Fabrys disease, familial hypercholesterolemia, polycystic
                                                                                                                                                                                                                                                                                                                   30-JUN-2000; 2000CN-0116935
                                                                                                                                                                                                                                                                                                                                              29-JUN-2001; 2001WO-CN01107
                                                                                                                                                                                                                                                                                                                                                                       14-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                  WO200211513-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABK12114 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                        (SHAN-)
                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EHD 9.57
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    ss; PCR; EHD 9.57; fetation abnormality; tumour; primer;
deficiency; neural system dysfunction disease;
metabolism disorder.
                                                                                                                                                                                                                                                                                         SHANGHAI BIOWINDOW GENE DEV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CCTGGCCAACATGGYGAAACCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CCTGGCGAACATGGTGAACCCC 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29
                                                                                                                                                               Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP;
 BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein PCR primer
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 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
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                                                                                                                                                             40pp; Chinese.
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10 C; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11 C; 6 G; 5 T; 1 other;
                                                                                                                                                                                                     and encoding
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ω
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 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18.4;
No. 2
                                                                                                                                                                                                     polynucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB
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                                                                                                                                                                                                     used
                                                                                                                                                                                                       in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29;
                                                                                                                                                                                                     diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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31

ACCACCCTGGCCAACATGGTGAA 9

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RESULT 13
ABK14174/c
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                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                      Query Match
                                                                                                                                             27605
                                                                         Human II aminoacyl-tRNA synthetase 9
                                                                                                                                                                                                                                                                          11.88 and the DNA sequence encoding the polypeptide. The DNA and protein are used in diagnosis and treatment of malignant tumour and dysmorphia of development of an embryo. This sequence represents a PCR primer used for cloning of cDNA which encodes the insulin like growth factor binding protein 11.88 of the invention.
                                                                                             31-AUG-2001
                                                                                                                                  AAH27605 standard;
                                                                                                                                                                                                                                                         Sequence 33
                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-172159/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unidentified
        Homo sapiens.
                        Human; II aminoacyl-tRNA synthetase 9; cytostatic; antiviral; immunomodulatory; antiinflammatory; cancer; haemopathy; human immunodeficiency virus; HIV; infection; immunological dinflammatory disease; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                The invention relates to an insulin like growth factor binding protein 11.88 and the DNA sequence encoding the polypeptide. The DNA and prote
                                                                                                                                                                                                                                                                                                                                                             Insulin like growth factor binding protein 11.88 and encoding polynucleotide, used in diagnosis and treatment of malignant tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200212493-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Insulin like growth factor binding protein 11.88; embryo development dysmorphia; malignant tumour; g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Insulin like growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABK14174;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABK14174 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                (BIOW-) BIOWINDOW GENE DEV INC SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-JUN-2000; 2000CN-0116491.
                                                                                                                                                                                                                                Local
                                                                                                                                                                                31
                                                                                                                                                                                          1 ATCAACCTGGCGAACATGGTGAA 23
                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                            4; Page 13; 38pp; Chinese.
                                                                                                                                                                                ACCAGCCTGGCCAACATGGTGAA 9
                                                                                                                                                                                                                                                        BP; 5 A; 11 C; 8 G; 9 T; 0 other;
                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2001WO-CN00951
                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first
                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 factor binding protein 11.88 cDNA PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
                                                                                                                                                                                                                            60.7%;
                                                                                                                                  34
                                                                                                                                  ВP
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                                                                                                                                                                                                                     0,
                                                                                                                                                                                                                              Score 18.2;
Pred. No. 2
                                                                                                                                                                                                                     Mismatches
                                    infection; immunological disease;
                                                                        PCR primer 4.
                                                                                                                                                                                                                              .9e+02;
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                                                                                                                                                                                                                                      24;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene therapy;
                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ss; cytostatic;
                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                   0;
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RESULT 15
AAA04312/c
.ID AAA043
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Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to an isolated polypeptide of human II aminoacyl-tRNA synthase 9. The polypeptide comprises the 86 amino acid sequence defined in the specification, or its fragment, analogue or derivative. The polypeptide and the polynucleotide encoding it are useful in the diagnosis and treatment of malignant tumours, haemopathy, HIV infection, immunological diseases and various inflammatory diseases. The polynucleotide is useful as a primer for nucleic acid amplification reactions or as a probe for hybridisation reactions, or in producing gene chips or microarrays. The polypeptide is useful for screening minics, agonists, antagonists or inhibitors, or for use in peptide fingerprinting identification. The present sequence is a primer which was used to amplify the polynucleotide encoding the polypeptide of the invention.
07-MAY-1998;
03-MAY-1999;
                                                                                                                                                                                                                                                                                                             Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis; polycystic kidney disease; von Willebrands disease; forensic; human; tuberous sclerosis; hereditary hemorrhagica telangiectasia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human II aminoacyl-tRNA synthase 9 and encoded polynucleotide, in diagnosis and treatment of malignant tumor, hemopathy, HIV infection, immunological diseases and various inflammation -
                                                                                                                                                                                                                                                                                                                                                                                  Polymorphism; hypertension; agammaglobulinemia; diabetes insipidus; Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAA04312 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 34 BP; 6 A; 11 C; 7 G; 10
                                                                        07-MAY-1999;
                                                                                                                                                                    EP955382-A2
                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                  Ehlers-Danlos
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Polymorphic fragment of hypertension associated gene GLUT4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA04312;
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The invention provides polymorphic fragments of genes associated with CC hypertension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic cc regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension of diagnostic assays. Where the polymorphisms have strong correlation with hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to find the precise role of a polymorphism in the disease, and this can be used to identify potential cc drugs which combat the diseases. The polymorphisms can be tested for association with other diseases e.g. agammaglobulinemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, Wiskott-Aldrich csyndrome, Fabrys disease, familial hypercholesterolemia, polycystic chidney disease, hereditary spherocytosis, von Willebrands disease, familial hypercholesterolemia, polycystic colonic polyposis, hereditary spherocytosis, von Willebrands disease, familial and colonic polyposis.
    Query Match
Best Local Similarity
Matches 21; Conserv
                                                                                                                                                                                                                                                                    colonic polyposis, Ehlers-Danlos syndrome, osteogenesis imperfecta, and acute intermittent porphyria. The polymorphic forms can also be used in forensics to identify individuals.
                                                                                                                                                                                     Sequence 29 BP; 5 A; 7 C; 7 G; 9 T; 1 other;
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(UYCA-) UNIV CASE WESTERN RESERVE.
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Search completed: June 21, 2003, 22:26:22 Job time: 165.531 secs

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26 ACCATTCTGGCYAACATGGTGAAACC 1

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US-09-09-913-171
US-08-256-368-14
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RESULT 2
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US-09-280-805-259; Sequence 259, Application US/09280805; Patent No. 6184212; GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pame APPLICANT: Graḥam, Brett P. Monia

Pamela

Nero,

NUMBER OF SEQUENCES: 2 CORRESPONDENCE ADDRESS:

OF INVENTION:

ANTISENSE MODULATION OF HUMAN MDM2 EXPRESSION

STREET: 66 Ea CITY: Marlton

66 East Main Street

Law Offices of Jane Massey Licata

ADDRESSEE:

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: OTHER INFORMATION: Synthetically generated oligonucleotides US-09-171-425-2
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APPLICANT: Schorr, Joachim
                                                                                                                                                                                                                                                                                                                                                                           Sequence 2, Application US/09171425A Patent No. 6465438
Query Match
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                                                                                                                                                                                                                                    TITLE OF INVENTION: NUCLEIC ACID VACCINATION FOR PARVOVIRAL INFECTIONS FILE REFERENCE: 08909/003001
CURRENT APPLICATION NUMBER: US/09/171,425A
CURRENT FILING DATE: 1998-10-19
                                                                                                                                                                                                     EARLIER APPLICATION NUMBER: PCT/EP97/01943
EARLIER FILING DATE: 1996-04-19
                                                                                                                                                                                                                                                                                                       APPLICANT: Baker, Henry J. APPLICANT: Smith, Bruce F.
                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 14
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ENFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
                                                                                     FEATURE:
                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                   TYPE: DNA
                                                                                                                                  ENGTH: 42
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TOPOLOGY: Lin
ANTI-SENSE: Yes
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REGERENCE/FOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
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APPLICATION NUMBER: 09/0
FILING DATE: March 26, 1
ATTORNEY/AGENT INFORMATION:
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Pred. No. 5.6e+02;
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15;
No.
DB 4;
7.6e+02;
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US-09-171-425-7/c
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US-09-171-425-6
                                                                   Query Match
Best Local Similarity
Matches 18; Conserv
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LENGTH: 48
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 7, Application US/09171425A Patent No. 6465438
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                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/171,425A CURRENT FILING DATE: 1998-10-19 EARLIER APPLICATION NUMBER: PCT/EP97/01943
                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Schorr, Joachim
APPLICANT: Baker, Henry J.
APPLICANT: Smith, Bruce F.
TITLE OF INVENTION: NUCLEIC ACID VACCINATION
FILE REFERENCE: 08909/003001
                                                                                                                                                                                                                                                               SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 14
                                                                                                                                                                                                                                                                                                  EARLIER FILING DATE: 1996-04-19
                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EARLIER APPLICATION NUMBER: PCT/EP97/01943
EARLIER FILING DATE: 1996-04-19
NUMBER OF SEQ ID NOS: 14
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CURRENT APPLICATION NUMBER: US/09/171,425A
CURRENT FILING DATE: 1998-10-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                      FEATURE:
OTHER INFORMATION: Synthetically generated oligonucleotides
                                                                                                                                                                                          TYPE: DNA ORGANISM: Artificial Sequence
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18; Conser
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                     1 ATCAACCTGGCGAACATGGTGAA 23
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ATCAACCTGGCTAAGAAGAAGAA 12
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                                                                      Conservative
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Pred. No.
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                                                                                                                                                                                                      Patent No. 6087485
GENERAL INFORMATION:
                                                                                                                                                                                                                                        Sequence 171, Application US/09009913
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LENGTH: 26 base pairs
TYPE: nucleic acid
COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                    APPLICANT: AxyS Pharmaceuticals, Inc.
TITLE OF INVENTION: Asthma Related Ge
NUMBER OF SEQUENCES: 339
CORRESPONDENCE ADDRESS:
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ATTORNEY/AGENT INFORMATION:
NAME: Highlander, Steven
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APPLICANT: Keller, Charles
APPLICANT: Akande, Olanike
TITLE OF INVENTION: GLUTATI
TITLE OF INVENTION: CANCER
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
SOFTWARE.
                                  COUNTRY: USA
ZIP: 94301
                                                                    STREET: 285 Ham:
CITY: Palo Alto
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                               Match 49.3%;
Local Similarity 88.9%;
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REGISTRATION NUMBER: 37,642
REFERENCE/DOCKET NUMBER: UTXC:492
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| 5 CCTGGTGGACATGGTGAA 22
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                                                                                                    285 Hamilton Ave, Suite 200
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Lopez-Berestein, Gabriel
Buolamwini, John
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                                                                                                                                                                      Asthma Related Genes
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RESULT 8
US-08-256-368-14/c
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Best Local Similarity 88.9%;
             TELEX: 248345
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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                                                                                                     APPLICATION NUMBER: US/08,
FILING DATE: 22-JUL-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30,3;
                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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SEQUENCE CHARACTERISTICS
                                                         TELEPHONE: (703)205-8000
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CORRESPONDENCE ADDRESS:
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                                            TELEFAX:
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                                                                                          REFERENCE/DOCKET NUMBER:
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Y: U.S.A.
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                                            (703)205-8050
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In Cells Of Spodoptera Frugiperda Infected With
Recombinant Baculovirus
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                                                                                                                                                                                    US/08/256,368
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. 8.7e+02;
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STRANDEDNESS:

single

nucleic acid

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В
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent No. 6451557
                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EARLIER APPLICATION NUMBER: PO3310
EARLIER FILING DATE: 1996-10-29
EARLIER APPLICATION NUMBER: PO4306
EARLIER FILING DATE: 1996-12-19
EARLIER APPLICATION NUMBER: PCT/AUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT FILING DATE: 1999-04-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: A METHOD FOR PRODUCING, TITLE OF INVENTION: HELICAL PROTEIN, AND YES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: RAMSHAW, Johr APPLICANT: WERKMEISTER,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EARLIER FILING DATE: 1997-10-29
NUMBER OF SEQ ID NOS: 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: METHOD (As Amended)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
                                                                                                                                                                                               quence 2, Appl
tent No. 53407
                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 41
                                       NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "PCR prime:
                                                                       APPLICANT: Wagner, Daniel B.
TITLE OF INVENTION: Assay Method Utilizing
TITLE OF INVENTION: Chemiluminescent Label
                                                                                                                    APPLICANT:
                                                                                                                                                               APPLICANT:
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            STREET:
CITY: Palo Alto
                              ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/09/297, 269
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                                                                                                                                                                                                           Application US/07718490
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               3401
                                                                                                                    Pease, John S.
Daniloff, Yuri
                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                   Kirakossian,
                                                                                                                                                                  Ullman,
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            Syntex (U.S.A.) Inc.
01 Hillview Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Paul R.
                                                                                                                                                                 Edwin F.
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                                                                                                                                                   Hrair
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Pred. No. 1
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Pred. No. 1

    Mismatches

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                                                                                      Photoactivated
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US-07-718-490-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
                                                                                                                                             STREET: 340.
STREET: DATA
CITY: Palo Alto
CTATE: California
" S.A.
                                         CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                   APPLICANT: Wagner, Daniel B.
TITLE OF INVENTION: ASSAY ME
TITLE OF INVENTION: Chemilum
NUMBER OF SEQUENCES: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FRAGMENT TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                    COMPUTER READABLE FORM:
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CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN: K12 DNAJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Leitereg, Theodore J. REGISTRATION NUMBER: 28,319
REFERENCE/DOCKET NUMBER: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 1: CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US
FILING DATE: 19910620
                                                          SOFTWARE:
                                                                                                    MEDIUM TYPE:
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                           APPLICATION NUMBER:
                                                                                                                                                                                                           ADDRESSEE:
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                                                                                                                                                                                                                                                                                                Daniloff, Yuri
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                                                                                                                                                                                                                                                                                                                             Kirakossian, Hrair
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                                                       PatentIn Release #1.0, Version #1.25
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415) 496-3529
                                                                                                    Floppy disk
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                                                                                                                                                                                         Hillview Avenue
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                                                                                                                                                                                                                                                                                                               John S.
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ATTORNEY/AGENT INFORMATION:

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RESULT 12
US-08-471-496-7
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Best Local Similarity
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                                                               INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                           ZIP: 20005-3934

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                         REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US95/02645
EILING DATE: 01-MAR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
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APPLICANT: CAO, LIANG
APPLICANT: ROSEN, CRA
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ORIGINAL SOURCE:
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                                              SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/471,496 FILING DATE: 06-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Leitereg, Theodore J. REGISTRATION NUMBER: 28,319
REFERENCE/DOCKET NUMBER: 27
                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM:
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TYPE: NUCLEIC ACID
STRANDEDNESS: single
STRANDEDNESS:
                                                                             TELEPHONE:
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            TYPE: hucleic acid
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5798223
                                                                                                                                                            STEFFE, ERIC K
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                                                                                                                                                                                                                                                                                                                                                                                                                 USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ROSEN, CRAIG
                                                                           (202) 371-2540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NO
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                                                                                            (202) 371-2600
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HUMAN AMINE TRANSPORTER
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Pred. No. 1.
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US-09-139-675-7
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            GENERAL INFORMATION:
APPLICANT: Li, Yi
APPLICANT: Cao, Liang
APPLICANT: Rosen, Craig
TITLE OF INVENTION: Human Amine Transporter
FILE REFERENCE: 1488.0830003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 7, Application US/08894840 Patent No. 5859200
                                                                                                    Sequence 7, Application US/09139675A Patent No. 6117426
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Best Local (
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CURRENT APPLICATION NUMBER: US/09/139,675A
                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (202) 371-254 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (202) 371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Relea
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/0
FILING DATE: '29-AUG-1997
                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: HUMAN AMINE TRANSPORTER NUMBER OF SEQUENCES: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                 Local Similarity 70.48; nes 19; Conservation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: LIVE CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 20005-3934
                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
TOPOLOGY: lir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: , Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX P.L.L.C. STREET: 1100 NEW YORK AVENUE, SUITE 600
                                                                                                                                                                                                                                                                                                                                                                                           LENGTH:
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CAO, LIANG
ROSEN, CRAIG A.
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No. 1.6e+03;
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Pred. No. 1
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SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 7

EARLIER FILING DATE: 1995-06-06 NUMBER OF SEQ ID NOS: 9

EARLIER APPLICATION NUMBER: WO PCT/US95/02645
EARLIER FILING DATE: 1995-03-01
EARLIER APPLICATION NUMBER: US 08/471,496

CURRENT FILING DATE: 1998-08-25

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TOPOLOGY:
US-08-853-774-11
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-139-675-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 70.4
19; Conservative
                                                                                                                                                                 Query Match
Best Local S
        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-853-774-11
                                                                                               INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 28 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     quence 11, Application US/08853774 (tent No. 6265557
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: PastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSEE: Knobbe, M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Nehlsen Cannarella, Sandra
APPLICANT: Fagoaga, Omar
APPLICANT: Szalay, Aladar
TITLE OF INVENTION: ABO HISTO-BLOOD GROUP O ALLELES OF THE BABOON
                                                                                                                                                                                                                                                                            FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: US
ZIP: 92660
                                                                                                                                                       TELEPHONE: 714/760-9503
                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Newport Beach
STATE: CA
                                                                                     STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                           USA
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Pred. No. 1.9e+03;
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Pred. No. 1.6e+03;
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8 TGGCGAACATGGTGAACCCCAT 29

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Search completed: June 22, 2003, 00:01:32 Job time : 36.625 secs

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Maximum Match
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Match Length
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Gapop 10.0 , Gapext 1.0
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1: /cgn2_6/ptodata/2/pubpna.

2: /cgn2_6/ptodata/2/pubpna.
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Copyright (c) 1993 - 2003 Compus
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US-09-860-670-233
US-10-085-906-77
US-09-530-139-69
US-09-530-139-70
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Sequence 11, Appl
Sequence 11, Appl
Sequence 233, Appl
Sequence 233, Appl
Sequence 69, Appl
Sequence 70, Appl
Sequence 35, Appl
Sequence 17, Appl
Sequence 259, Appl
Sequence 32, Appl
Sequence 6391, Appl
Sequence 6491, Appl
Sequence 4044, Appl
Sequence 36, Appl
Sequence 37, Appl
Sequence 36, Appl
Sequence 37, Appl
Sequence 36, Appl
Sequence 37, Appl
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13.6	13.6	13.6	13.6	13.6	13.6	13.6	13.6	13.6	13.8	13.8	13.8	13.8	13.8	14	14	14	14.2	14.2	14.2	14.2	14.2	14.2	14.2	14.4	14.6
45.3	45.3	45.3	45.3	45.3	45.3	45.3	45.3	45.3	46.0	46.0	46.0	46.0	46.0	46.7	46.7	46.7	47.3	47.3	47.3	47.3	47.3	47.3	47.3	48.0	48.7
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US-U9-8//-4/8-4115	US-09-780-533A-4376	US-09-730-289B-2710	US-10-155-233-10	US-09-801-274-1365	US-09-864-636A-1614	US-10-085-906-44	US-09-529-063-86	US-10-085-906-365	US-10-146-938-1	US-09-930-423-2860	US-09-930-423-2785	US-10-192-926-18	US-10-025-201-13	US-09-832-292-24	US-10-098-263B-25783	US-09-940-185-2688	US-09-930-423-2915	US-09-877-478-4122	US-09-780-533A-4387	US-09-780-533A-4302	US-09-987-025-12	US-09-924-125-4	US-09-964-277-15	US-09-755-665-108	US-10-085-906-241
Sequence 4115, Ap	Sequence 4376, Ap	Sequence 2710, Ap	Sequence 10, Appl	Sequence 1365, Ap	Sequence 1614, Ap	Sequence 44, Appl	Sequence 86, Appl	Sequence 365, App	Sequence 1, Appli				Sequence 13, Appl		Sequence 25783, A	Sequence 2688, Ap		Sequence 4122, Ap	Sequence 4387, Ap	Sequence 4302, Ap	Sequence 12, Appl	Sequence 4, Appli	Sequence 15, Appl	ς.	Sequence 241, App

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## ALIGNMENTS

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RESULT 1
US-09-964-667-11
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                                                                                                                                INFORMATION FOR SEQ ID NO: 11: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 20005-3934

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF CONTROL OF THE CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
Sterne, Kessler, Goldstein
ADDRESSEE: Sterne, Kessler, Gulte 600
                                                                                                                                                                                                 NAME: Esmond, Robert W.
REGISTRATION NUMBER: 32,893
REFERENCE/DOCKET NUMBER: 061
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/964,667
FILING DATE: 28-Sep-2001
CLASSIFICATION: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: cDNA
SEQUENCE DESCRIPTION: SEQ ID NO: 11:
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Wands, Jack R
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                                                              LENGTH: 30 base pairs TYPE: nucleic acid STRANDEDNESS: single
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                                                                                                                                                                                 TELEFAX: 202-371-2540
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Treatment
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of Alzheimer's Disease

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US-09-964-666-11
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                                                         Seguence 11, Application US/09964412
Patent No. US20020129391A1
GENERAL INFORMATION:
APPLICANT: de la Monte, Suzani
                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
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TITLE OF INVENTION: Transgenic Animals and Cell Lines for Screening Drugs Effective for the Treatment or Prevention
                                                                                                                                                                                                                                                                                                                                                   SEQUENCE DESCRIPTION: SEQ ID NO: 11:
                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
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FILING DATE: 28-Sep-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single
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TELEFAX: 202-371-2540
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                                        de la Monte, Suzanne
Wands, Jack R.
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Wands, Jack R.
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                                                                                                                                                                                                                                                                                100.0%;
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Pred. No. 0.00063;
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RESULT 4
US-09-860-670-233
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                                                                                                                                                                 Prior application data removed - consult PALM or file wrapper NUMBER OF SEQ ID NOS: 289
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 233
LENGTH: 49
TYPE: DNA
ORGANISM: Homo sapiens
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Best Local Similarity
                                                                        Matches
                                                                                            Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 233,
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                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies FILE REFERENCE: PA127P1 CURRENT APPLICATION NUMBER: US/09/860,670 CURRENT FILING DATE: 2001-05-21
                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Ruben et al.
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TELEFAX: 202-371-2540
INFORMATION FOR SEQ ID NO: 11:
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APPLICATION NUMBER: US/09/964,412
ETITUT CAMPS 2025-2027
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox, P.L.L.C.
STREET: 1100 New York Ave., Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE DESCRIPTION: SEQ ID NO: 11:
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16 ACCATCCTGGCTAAAATGGTGAAACCC 42
                 -1 ATCAACCTGGCGAACATGGTGAACCCC 27
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                                                                                          Similarity
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TYPE: nucleic acid
STRANDEDNESS: single
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                                                                        Conservative
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                                                                                          63.3%;
81.5%;
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                                                                                            Score 19;
Pred. No.
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Pred. No. 0.00063;
                                                                        Mismatches
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                                                                                                            DB 9;
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RESULT 7
US-09-530-139:70
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; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-09-530-139-69
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                                                                                                                                                                  Query Match
Best Local Similarity
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Best Local Similarity
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CURRENT FILING DATE: 2002-02-27
PRIOR APPLICATION NUMBER: US 60/126,215
PRIOR FILING DATE: 1999-03-25
PRIOR APPLICATION NUMBER: US 09/534,061
PRIOR FILING DATE: 2000-03-24
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TITLE OF INVENTION: COSTIMULATORY RECEPTOR LOCUS AND USES THEREOF
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                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: EP 97308538.4
PRIOR FILING DATE: 1997-10-27
NUMBER OF SEQ ID NOS: 77
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: VAN DER LOGT, CORNELIS PAUL
TITLE OF INVENTION: MULTIVALENT ANTIGEN-BINDING PROTEINS
FILE REFERENCE: 60113/268075/ASH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: PCT/US00/07938 PRIOR FILING DATE: 2000-03-24
                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: PCT/EP98/0699: PRIOR FILING DATE: 1998-10-27
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CURRENT FILING DATE: 2000-04-27
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                                                                                                                                                                                                                                                                                                      TYPE: DNA
                                                                                                                                                                                                                                                                                                                        LENGTH: 29
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                                                                          ACCTGGTGAACCTGGAGAACCAC 4
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No. US20030092892A1
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Pred. No. 3.7e+02;
0; Mismatches 2;
                                                                                                                                                                  Score 16.6; DB 9;
Pred. No. 4.5e+02;
                                                                                                                                                   Mismatches
                                                                                                                                                                                    DB 9;
                                                                                                                                                                                  Length 29
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US-09-885-441-35/c
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                                                                                                                                                                                                                                      SEQ ID NO 35
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                                                                              Matches
                                                                                              Query Match
Best Local
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Best Local Similarity 85.7%;
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TITLE OF INVENTION: MULTIVALENT ANTIGEN-BINDING PROTEINS
FILE REFERENCE: 60113/268075/ASH
CURRENT APPLICATION NUMBER: US/09/530,139
CURRENT FILING DATE: 2000-04-27
PRIOR APPLICATION NUMBER: PCT/EP98/06991
PRIOR PILING DATE: 1998-10-27
PRIOR PILING DATE: 1998-10-27
PRIOR APPLICATION NUMBER: EP 97308538.4
PRIOR APPLICATION NUMBER: EP 97308538.4
NUMBER OF SEQ ID NOS: 77
                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 60/212,844 PRIOR FILING DATE: 2000-06-21 PRIOR APPLICATION NUMBER: US 60/244,171 PRIOR FILING DATE: 2000-10-31 PRIOR APPLICATION NUMBER: US 60/279,766 PRIOR FILING DATE: 2001-03-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Regulation of Human Eosinophil Serine TITLE OF INVENTION: Protease-1-Like Enzyme FILE REFERENCE: 04974.00512
                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: PCT/PRIOR FILING DATE: 2001-06-20 NUMBER OF SEQ ID NOS: 58
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                                                                                                                                                                                               LENGTH: 24
TYPE: DNA
                                                                                                                                                                                                                                                        SOFTWARE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/885,441
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CTGGTCAACATGGTGAAACCC
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                                                                              Conservative
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                                                                                              54.0%;
85.7%;
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                                                                                              Score 16.2; DB 10
Pred. No. 6.5e+02;
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Pred. No. 6.5e+02;
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RESULT 9
US-10-123-170-17
; Sequence 17, Application US/10123170

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US-09-752-983-259
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Patent No. US2001
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Best Local
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CURRENT APPLICATION NUMBER: US/10/123,170
CURRENT FILING DATE: 2002-04-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: RECOMBINANT SEGMENTED NEGATIVE STRAND VIRUS CONTAINING BICISTRON TITLE OF INVENTION: SEGMENT WITH A DUPLICATION OF ITS 3' NONCODING FLANKING SEQUENC TITLE OF INVENTION: AND THERAPPUTIC COMPOSITIONS CONTAINING THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT FILING DATE: 2002-04-17
PRIOR APPLICATION NUMBER: 60/283,957
PRIOR FILING DATE: 2001-04-17
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                            TELEFAX: 609-810-1454
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
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                                                                                  REFERENCE/DOCKET NUMBER: IS TELECOMMUNICATION INFORMATION: TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                            SOFTWARE: WORDPERFECT 6.0 CURRENT APPLICATION DATA:
                                                                                                                                                                APPLICATION NUMBER: 09/28
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: DISKETT
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                                                                                                                                       NAME: Licata, Jane Massey REGISTRATION NUMBER: 32,2
                                                                                                                                                                                                                                                              APPLICATION NUMBER: FILING DATE: 02-Jar
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nes 19; Conserv
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Nucleic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       66 East Main Street
                  20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/09752983
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                                                                                                                                                                                                                                                                                                                                                IBM PC
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79.2%;
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Pred. No. 8.
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8.3e+02;
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; ORGANISM: Homo sapiens US-09-801-274-1110
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US-09-800-631-32/c
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Best Local (
                                                                                                                           SEQ ID NO 1110
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                                                                                                                                                     PRIOR APPLICATION NUMBER: US 60/206,129
PRIOR FILING DATE: 2000-05-22
NUMBER OF SEQ ID NOS: 1802
                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/801,274
CURRENT FILLING DATE: 2001-03-07
PRIOR APPLICATION NUMBER: US 60/187,510
PRIOR FILING DATE: 2000-03-07
                                                                                                                                                                                                                                                                 APPLICANT: Ireland, James S.
APPLICANT: Lander, Eric S.
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE
FILE REFERENCE: 2825.2009-001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Hong Zhang
APPLICANT: Jacqueline Wyatt
TITLE OF INVENTION: ANTISENSE MODULATION OF BH3 INTERACTING DOMAIN DEATH AGONIST E
FILE REFERENCE: ISPH-0544
                                                                                                                                            SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                APPLICANT: Cargill, Michele
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TYPE: DNA
                                                                                              TYPE: DNA
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TOPOLOGY: Li
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85.0%;
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Pred. No. 1.8e
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Pred. No. 1.7e+03;
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 07/882,712 PRIOR FILING DATE: 1992-05-14
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PRIOR FILING DATE: 2000-03-20
PRIOR APPLICATION NUMBER: US 09/636,385
PRIOR FILING DATE: 2000-08-09
                                                                                                                                                                           OTHER INFORMATION: n stands for inverted
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OTHER INFORMATION: 2'-0-Methyl
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OTHER INFORMATION: Phosphorothioate
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OTHER INFORMATION: Phosphorothioate
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APPLICATION NUMBER: US
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Blatt, Larry
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Morrissey, Dave
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Pred. No. 1
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RESULT 14

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CURRENT APPLICATION NUMBER: US/10/098,263B
CURRENT FILING DATE: 2003-01-08
PRIOR APPLICATION NUMBER: 60/276,759
PRIOR FILING DATE: 2001-03-16
NUMBER OF SEQ ID NOS: 131066
SOFTWARE: Microarray Probe Sequence Listing G
SEQ ID NO 8701
LENGTH: 25
TYPE: DNA
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US-10-098-263B-8701
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Matches 18; Conserv
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APPLICANT: McSwiggen, Jim
APPLICATION: Method and Reagent for Inhibiting Hepatitis B Virus
FILE REFERENCE: MBHB00-845-H (400/029)
FILE REFERENCE: MBHB00-845-H (400/029)
CURRENT APPLICATION NUMBER: US/09/877,478
CURRENT APPLICATION NUMBER: 2001-12-31
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APPLICANT: Draper, Kenneth
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
                                                                                                                                                                                                                                                                                                            APPLICANT: Mittman, Michael
TITLE OF INVENTION: Human Microarray
FILE REFERENCE: 3118.1
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PRIOR FILING DATE: 2000-10-24
PRIOR APPLICATION NUMBER: US 08/193,627
PRIOR FILING DATE: 1994-02-07
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                                                                                                       ORGANISM: Homo sapien
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APPLICATION NUMBER: US 08/434,504
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Search completed: June 22, 2003, 03:18:19
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Contact: Yutaka Suzuki
Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki, Y., Yoshitomo-Nakagawa, K., Maruyama, K., Suyama, A. and Sugano, S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
Location/Qualifiers
                                                                                                                 Y., Nakamura,Y., Suyama,A. and Sug
Diverse transcriptional initiation
mapping of mRNA start sites
EMBO Rep. 2 (5), 388-393 (2001)
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R05734
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ion revealed
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AU1042536 AU104233

AU104439 AU1042521

AU102521 AU102528

AU103129 AU105233

AU103129 AU105233

AU105233 AU105233

AU105233 AU105233

AU105742 AU105742

R91494 yq08f12.s1

AA912807 0143a05.s

AU102522 AU102523

AU104028 AU1042531

AU104028 AU104028

AU1040437 AU104437

AU309847 1W001708

H82905 yq46e11.r1

H88764 yw23f12.r1

C20880 HUMCS000494

AU10244156 AU244156

N57213 yw65b02.r1
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H56911 yr07d07.s1
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RESULT 3
N80349/c
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Contact: Yutaka Suzuki
Contact: Yutaka Suzuki
Department of Virology
Institute of Medical Science, University of Tokyo
Institute of Medical Science, University of Tokyo
N80349 33 bp mRNA linear EST 02-APR-1996 zal3c12.rl Soares fetal liver spleen lNFLS Homo sapiens CDNA clone IMAGE:292438 5' similar to gb|Ll3704|HUMSCALUB Human scrNA molecule, transcribed from Alu (rRNA); gb:X53795_rnal INDUCIBLE MEMBRANE
                                                                                                                                       1 ATCAACCTGGCGAACATGGTGAACCC
                                                                                                                                                                                                                                                                                                                                                                                           Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano,S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 50)
Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J., Sese,J., Hat, Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo,K., Sakaki,Y., Nakamura,Y., Suyama,A. and Sugano,S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mapping of mRNA start sites EMBO Rep. 2 (5), 388-393 (20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Diverse transcriptional initiation revealed by fine, large-scale
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/note="Differential display comparison of untreated
dimethylfumarate treated U937 cells"
a 10 c 12 g 14 t
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                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="LNG05691"
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/note="Differential display comparison of untreated dimethylfumarate treated U937 cells"
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/clone="COL01940"
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Pred. No. 1
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Pred. No. 75;
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AUTHORS
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AU102526/c
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Best Local :
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33 CATCCTGGCTAACATGGTGAANCCC
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Endammalla; Eutheria; Primates; Catarrhini; Hominidae; I (bases 1 to 50)
Suzuki, Y., Taira, H., Tsunoda, T., Mizushima-Sugano, J., H., Ota, T., Isogai, T., Tanaka, T., Morishita, S., Okub, Y., Nakamura, Y., Suyama, A. and Sugano, S.
                                                                                                                                                                                                                        AU102526 Sugano Homo sap
COLF1341, mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (1995)
                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: est@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4444 Forest Park Parkway, Box 8501, St. Louis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Washington University School of Medicine
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314 286 1810
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                                                                                                                                                                                                                                             Sugano Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (Pharmacia), digested with Pac I and cloned into the Pac I and Eco RI sites of the modified pT7T3 vector. Library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /dev_stage="20 week-post conception fetus"
/lab_host="DH10B (ampicillin resistant)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /db_xref="taxon:9606"
/clone="IMAGE:292438"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone_lib="Soares fetal.liver spleen lNFLS
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Pred. No. 1
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RESULT 5
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Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
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Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano,S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
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21270072
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Department of Virology
Institute of Medical Science, University of Tokyo 4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Jar Email: ysuzuki@ims.u-tokyo.ac.jp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J., Sese,J., Ha, Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo,K., Sakaki,Y., Nakamura,Y., Suyama,A. and Sugano,S.
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AU105702 Sugano Homo sapiens
HRC11393, mRNA sequence.
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Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Jap
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                                                                                                                /Clone_lib="Sugano Homo sapiens cDNA library"
/note="Differential display comparison of unt
dimethylfumarate treated U937 cells"
a 16 c 12 g 9 t
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/note="Differential display comparison of untreated
dimethylfumarate treated U937 cells"
/note="Differential display comparison of untreated
dimethylfumarate treated U937 cells"
/note="Differential display comparison of untreated
dimethylfumarate treated U937 cells"
/note="Differential display comparison of untreated updates"
/note="Diff
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/db_xref="taxon:9606"
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/db_xref="taxon:9606"
                                                                                                                                                                                                                                         /clone="HRC11393"
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Pred. No. 1.9e
0; Mismatches
   Score
Pred.
   20.6; DB 9;
No. 1.9e+02;
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14 ACCAGCCTGGCCAACATGGTGAAACCC
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Tissue Procurement: David G. Bostwick, M.D., Rodr
M.D., Michael R. Emmert-Buck, M.D., Ph.D.
cDNA Library Preparation: David B. Krizman, Ph.E
cDNA Library Arrayed by: Greg Lennon, Ph.D:
DNA Sequencing by: Washington University Genome
Clone distribution: MCI-CGAP clone distribution
found through the I.M.A.G.E. Consortium/LLNL at:
                   AU103189
AU103189 Sugano Homo sap
HRC04023, mRNA sequence.
AU103189
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44 bp mRNA linear EST 19-A ng79a04.sl NCI_CGAP_Pr6 Homo sapiens cDNA clone IMAGE:940974 similar to gb:M91159 !!!! ALU CLASS E WARNING ENTRY !!!! (HU
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Contact: Robert Strausberg, Ph.D.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae
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National Cancer Institute, Cancer Genome Anatomy
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                                                      Sugano Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                       /note="Vector: pAMP10; mRNA made from prostatic intraepithelial neoplasia.(low-grade), cDNA made boligo-dT priming. Non-directionally cloned. Size-selected on agarose gel, average insert size Reference: Krizman et al. (1996) Cancer Research
                                                                                                                                                                                                                                                                                                                                        56:5380-5383."
14 c
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/clone="IMAGE:940974"
/clone_lib="NCI_CGAP_Pr6"
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/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
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 GI:13552710
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Pred. No. 2.2e
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano
,S. Construction and characterization of a full length-enriched and
a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
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Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J., Sese,J., Hat,H., Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo,K., Sakaki,Y., Nakamura,Y., Suyama,A. and Sugano,S.
Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
EMBO Rep. 2 (5), 388-393 (2001)
  Source:
                                                                                 Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                      Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. Louis,
                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 33)
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                   High qality sequence starts: 1 High qality sequence stops: 1
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Contact: Wilson RK
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Department of Virology
Institute of Medical So
                                                              Email: est@watson.wustl.edu
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/note="Differential display comparison of untreated
dimethy.lfumarate treated U937 cells"
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/db_xref="taxon:9606"
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Minatoku, Tokyo 108-8639, Jap
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Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Su,H., Ota,T., Isogai,T., Tanaka,T., Morishita,Y., Nakamura,Y., Suyama,A. and Sugano,S.
Diverse transcriptional initiation revealed b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AU102524 Sugano Homo sar
COL07268, mRNA sequence
22;
                                                                                                                                                                                                             Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A.
,S. Construction and characterization of a full length-er
a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (
                                                                                                                                                                                                                                                         Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Jap
Email: ysuzuki@ims.u-tokyo.ac.jp
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/clone_lib="Sugano Homo sapiens cDNA library"
/note="Differential display comparison of untreated and dimethylfumarate treated U937 cells"
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/db_xref="GDB:489723"
/db_xref="taxon:9606"
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COL08447, mRNA sequence.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 50)
Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J., Sese,J., Hat,H., Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo,K., Sakaki,Y., Nakamura,Y., Suyama,A. and Sugano,S.
Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
EMBO Rep. 2 (5), 388-393 (2001)
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Contact: Yutaka Suzuki
Department of Virology
Institute of Medical Science, University of Tokyo
A-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano,S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
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                                                                                                                                                                                                    HRC07143, mRNA sequence.
AU102529
AU102529.1 GI:13552049
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Diverse transcriptional initiation revealed by fine, large-scale
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EMBO Rep. 2 (5), 388-393 (2001)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/db_xref="taxon:9606"
/clone="CDL08447"
/clone_lib="Sugano Homo sapiens cDNA library"
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Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano
,S. Construction and characterization of a full length-enriched and
a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997)...
Location/Qualifiers
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AU102530 Sugano Homo sapiens
HSI02245, mRNA sequence.
    Similarity.
22; Conserv
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Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano
,S. Construction and characterization of a full length-enriched and
a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
                                                                                                                                                                                                                                                                                                                                                                                         Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Jap
Email: ysuzuki@ims.u-tokyo.ac.jp
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Institute of Medical Science, 4-6-1, Shirokanedai, Minatoku,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Yutaka Suzuki
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/note="Differential display comparison of unt
dimethylfumarate treated U937 cells"
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dimethylfumarate treated 0937 cells"
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Score 19; DB 9; Le
Pred. No. 8.3e+02;
D; Mismatches 5;
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                          Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J.,
H., Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo
,Y., Nakamura,Y., Suyama,A. and Sugano,S.
Diverse transcriptional initiation revealed by fine, I
mapping of mRNA start sites
EMBO Rep. 2 (5), 388-393 (2001)
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   Contact:
                                                                                                                      Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 50)
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Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email ysuzuki@ims.u-tokyo.ac.jp
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Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
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1 (bases 1 to 50)
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/note="Differential display comparison of untreated and dimethylfumarate treated U937 cells"
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   Yutaka Suzuki
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/db_xref="taxon:9606"
/clone="NBLAN362NF"
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Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano
y.S. Construction and characterization of a full length-enriched and
a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
                                                                                                                                                                                                                                                                                                                                                                                    Suzuki, r., Yoshitomo-Nakagawa, R., Maruyama, K., Suyama, A. and Sugano, S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Yutaka Suzuki
Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites EMBO Rep. 2 (5), 388-393 (2001) 21270072
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J., ,H., Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo,Y., Nakamura,Y., Suyama,A. and Sugano,S.
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AU103194 Sugano Homo sapiens CDNA
NBLAN2977NF, mRNA sequence.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/clone="Differential display comparison of untreated dimethylfumarate treated U937 cells"
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/note="Differential display comparison of unt
dimethylfumarate treated U937 cells"
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SUMMARIES

## ACCESSION VERSION KEYWORDS SOURCE RESULT 1 AR071993/c LOCUS DEFINITION REFERENCE Result a 00000 000 000 000000000 00000 C AUTHORS TITLE ORGANISM No Score 111.4 Sequence 23 AR071993 AR071993.1 1 (bases 1 to 25) Watson, J.D. and Rudert, F. CD95 regulatory gene sequences CD95 regulatory gene 315-JUN-1999; Patent: US 5912168-A 215-JUN-1999; Unknown. Unclassified Unknown AR071993 Match Query 23 Leng 25 bp from patent US 5912168 GI:7222881 DΒ AR110255 AX467734 AX462456 BD013007 BD013007 PFDFRGMENY A07187 AR077392 AR077493 AR118012 AR158850 AR158851 AR158852 AR158853 AR158854 AR173023 AR000120 AR158847 AR158848 AR158849 AX294160 AX296756 AX289527 AX292123 AR117988 I95709 AR071993 I06372 AR193144 AR154129 AX378274 AR049979 AR083596 AR182515 AR077405 AR077368 ID 107406 ALIGNMENTS DNA linear AR077392 Sequence AR077493 Sequence AR118012 Sequence 195733 Sequence 61 AX042456 Sequence BBD13007 Novel G p BBO110040 Novel G p L08009 Bacteriopha A07187 primer 8/1 L08009 Bacteriopha A0718 primer 8/1 E59618 Human prote 105106 Sequence 5 107406 Sequence 18 107637 Sequence 6 171445 Sequence 15 AR158850 AR158851 AR158853 AR158853 AR158854 AR173023 AR000120 AR205963 AR000120 AR205963 AR0001255 AR467734 AX042456 AX296756 AX289527 I95733 Sequence 6 AR077636 Sequence AX201429 Sequence AR1 AR1 AR1 AR077405 Sequence AR117988 Sequence I95709 Sequence 37 AR193144 Sequence AR154129 Sequence AR071993 Sequence 106372 Sequence 33 AR049979 AX378274 Description PAT 1756 sequence 123 sequence 123 sequence 1274 sequence 1374 sequence 1375 sequence 1375 sequence 13848 sequence 13848 sequence 13851 sequence 13852 sequence 13853 sequence 13854 sequence 13854 sequence 13854 sequence Sequence Sequence 18-FEB-2000 Sequence Sequence

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Mulvihill, E.R., Nexo, B.A., Yoshitake, S.,
Hashimoto, A. and Yuzuriha, T.
Mutant t-PA with kringle replacement
Mutant t-PA with kringle replacement
Patent: EP 0293934-Al 33 07-DEC-1988;
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106372
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Dean, N.M. and Cowsert, L.M.
Antisense inhibition of HPK/GCK-like kinase expression
Patent: US 6346416-A 29 12-FEB-2002;
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7 c 4 g
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AR154129
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AR077405
1 (bases 1 to 36)
Griffiths,A.David., William
Nissim,A., Winter,G.Paul.,
                                                                                                                                                                                                                                                                                                                                                                    1 (bases 1 to 36)
Griffiths, A. David., Williams, S. Cameron., Water
Nissim, A., Winter, G. Paul., Johnson, K. Stuart.
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12; Conserv
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12; Conserv
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Yeh,H.R. and Wick,C.H.
Detector for nucleic acid typing and methods of using the same
Patent: US 6238866.A.7. 29-MAY-2001;
                                     Unclassified
                                                                                                                                                                                                        1 CACTGCACTTNCCA 14
                                                                                                                                                                                                                                                                                                                                        Methods for producing recombinant vectors Patent: US 5962255-A 83 05-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                               Smith, A. John. Hammond
                                                                                                                                                                                                                                                                                                                                                                                                            Unclassified
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             Williams, S. Cameron.,
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  Johnson, K. Stuart.
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AR117988
  l Similarity
12; Conser
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Johnson,K.Stuart., Winter,G.Paul., Griffiths,A.David.,
Smith,A.John.Hammond. and Waterhouse,P.Michael.
Methods for producing members of specific binding pairs
Patent: US 5733743-A 37 31-MAR-1998;
Location/Qualifiers
                                                                                                                                                                                                                                                                                       1 (bases 1 to 36)
Johnson, K. Stuart., Winter, G. Paul., Griffiths, A. David., Smith, A. John. Hammond. and Waterhouse, P. Michael.
Methods for producing members of specific binding pair.
Patent: US 6140471-A 37 31-OCT-2000;
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/db_xref="taxon:32630"
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/db_xref="taxon:32630"
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RESULT 11 AX289527/c

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AX292123
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/db_xref="taxon:32630"
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1 (bases 1 to 18)
Borden, Pann. and Heller, R. Anand.
Borden, Pann and Heller, R. Anand.
Methods for measuring stromelysin-1 promoter activity
Patent: US 6338944-A 2 15-JAN-2002;
                                                                                                                                                                                                                                                                                                                                       1 CACTGCACTTNCC 13
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Borden, P.Ann. and Heller, R. Anand.
Human stromelysin-1 promoter
Patent: US 5824794-A 2 20-OCT-1998;
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Sequence 2 from patent
AR049979
AR049979.1 GI:5971971
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Obesity associated biallelic marker maps
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae
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Query Match 74.3%; Score 10.4; DB 6; Length 18;
Best Local Similarity 84.6%; Pred. No. 8.9e+04;
Matches 11; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Oy 1 CACTGCACTINCC 13
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Db 3 CACTGCCCTTACC 15

Search completed: June 21, 2003, 22:14:35

Job time: 284.094 secs

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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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1: /SIDS2/gcgdata/geneseg/genesegn-emb1/NA1980.DAT:*

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4: /SIDS2/gcgdata/geneseg/genesegn-emb1/NA1983.DAT:*

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410.621 Million cell updates/sec
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	9:	/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1988.DAT:*
	10:	/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1989.DAT:*
	11:	/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1990.DAT:*
	12:	/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1991.DAT:*
	13:	/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1992.DAT:*
	14:	/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1993.DAT:*
	15:	/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1994.DAT:*
	16:	/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1995.DAT:*
	17:	/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1996.DAT:*
	18:	/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1997.DAT:*
	19:	/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1998.DAT:*
	20:	/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1999.DAT:*
	21:	/SIDS2/gcgdata/geneseq/geneseqn-embl/NA2000.DAT:*
	22:	/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA2001A.DAT:*
	23:	/SIDS2/gcgdata/geneseq/geneseqn-embl/NA2001B.DAT:*
	24:	/SIDS2/gcgdata/geneseq/geneseqn-embl/NA2002.DAT:*
ed. No.	is	ed. No. is the number of results predicted by chance to have a
ore gre	ater	ore greater than or equal to the score of the result being printed,
d is de	rive	d is derived by analysis of the total score distribution.

Result		%				
NO.	Score	Match	Match Length DB	DB	ID.	Description
c 1	13	92.9	25	19	AAV23155	Primer -1299/-1/CA
c 2	13	92.9	27	20	AAX84455	PCR primer for BCN
ω	11.4	81.4	20	24	ABK44429	Human HPK/GCK-like
4	11.4	81.4	22	22	AAS03611	PCR primer rend9b1
ر ت	11.4	. 81.4	27	9	AAN81248	Probe 0-YNH24 to i
6	11.4	81.4	27	9	AAN82045	Probe O-YNH24 for
7	11.4	81.4	31	22	AAS04026	VNTR hybridisation
8	11.4	81.4	36	14	AAQ49016	Multimeric (SBP) a
9	11.4	81.4	41	22	AAH46243	Human neuronal thr

æ

PCR primer RT-R6 f	AAV45623	20	22		10.4	45	C	
Primer RT-R6 for d	AAZ28746	20	22		0	44	O	
Nicotiana benthami	AAD35625	24	21		1.0.4	43	O	
rimer used t	AAH25541	22	21		0	42	O	
Human GPRv21 DNA P	AAH73538	22	21		0	41		
Zsig66 sequencing	AAA49891	21	21	٠	0	40	a	
ulpha	AAT17610	17	21		0	39		
ĸ	AAT34465	17	21	74.3	10.4	38		
interleuki	ABK14343	24	20		0	37		
$^{\circ}$	AAH63063	22	20		0	36		
Oligonucleotide hy	AAH80512	22	20		0	ა 5	C	
leotide	AAH80511	22	20		0	34	O	
leotide	AAH80510	22	20		10.4	ω ω	C	
leotide	AAH80509	22	20		10.4	32	O	
leotide	AAH80508	22	20		10.4	31	C	
leotide	AAH80507	22	20		10.4	30	c	
Oligonucleotide hy	AAH80506	22	20		10.4	29	c	
nucleotide	AAH80505	22	20		0	28	c	
	AAS10648	22	20		0	27	c	
r A (Group	AAQ95921	16	20		0	26	O	
stromelys	ABA92015	24	18		0	25		
$\overline{}$	AAZ73123	21	18		0	24		
sequ	AAV62716	19	18	•	0	23		
Human yes1 biallel	AAX90213	20	17	•	0	22		
Oligonucleotide SE	ABH25833	23	13	•	0	21		
Oligonucleotide SE	ABH25832	23	13	٠.	10.4	20	ი	
. Human biallelic po	AAX06147	20	31		0	19		
ശ	ABK40815	24	47		11	18	O	
AI2 PCR p	AAA71469	21	28	•	11	17	C	
e oligonuc	ABI90167	24	24	•	11	16		
oligonuc	ABI90166	24	24	78.6	11	15	C	
oligonuc	97	24	24		11	14		
oligonuc	197	24	24		11	13	c	
oligonuc	1967	24	20		11	12	ი	
ture oligonucl	420	24	20	•	11	11	O	
Dog genomic marker	AAA66165	21	20		11	10		

## ALIGNMENTS

ВP

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RESULT 1
AAV23155/c
AAV23155;
XX
AC AAV23155;
XX
CD AAV23155;
DT 23-JUL-1998 (first entry)
XX
DE Primer -1299/-1/CAT for human
XX
CD95; regulatory region; sile
XW
CD95; regulatory regions for human
XX
DE
XY
OS Homo sapiens.
XX
PN
WO9808965-A2.
XX
P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CD95; regulatory region; silencer; enhancer; apoptosis; inhibitor; cancer; viral infection; neurodegeneration; autoimmune disease; human; gene therapy; transcription factor; PCR primer; ss.
                  New regulatory regions from the CD95 gene and transcription factors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Primer -1299/-1/CAT for human CD95 gene.
                                                                                                                                                                                                                                                                                                                                                                                (GENE-) GENESIS RES & DEV CORP LTD
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                     This sequence is a PCR primer for DNA encoding the brain cyclic nucleotide-gated ion channel (BCNG) of the invention. BCNG and BCNG-related proteins are useful in screening for compounds that modulate, interact or affect expression. Compounds, e.g. antagon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BCNG; brain cyclic nucleotide gated ion channel; epilepsy; hyperalgesia; Alzheimer's Disease; Parkinson's Disease; long QT syndrome; dyslexia; sick sinus syndrome; age-related memory loss; cystic fibrosis; sudden death syndrome; pacemaker rhythm dysfunction; sensory disorder; auditory disorder; respiratory disorder; attention deficit disorder; learning disability; drug addiction; therapy; PCR primer; ss.
                                                                                                                                                               Claim 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        autoimmune disease, e.g. by gene therapy for expressing transcription factors or expression of antisense sequences to inhibit transcription factor production. The regulatory nucleic acids and transcription factors are also useful for studying regulation of CD95 in vitro or in vivo, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This sequence is a primer for the CD95 gene, and is used to isolate the regulatory nucleic acids of the invention. The regulatory regions (silencers or enhancers) are involved in apoptosis, e.g. inhibition of CD95 expression will inhibit apoptosis. Regulation of apoptosis is useful in treatment of cancer, (retro)viral infection, neurodegeneration and partitions of decorate the contraction of the contraction o
                                                                                                                                                                                                                                                                                         WPI; 1999-418922/35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-MAY-1998;
23-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAX84455 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            that interact with them - treatment of cancer, vira
     agonists,
                                                                                                                                                                                                            nucleotide-gated
                                                                                                                                                                                                                                    An isolated nucleic acid encoding a brain
                                                                                                                                                                                                                                                                                                                                                                   Bartsch D,
                                                                                                                                                                                                                                                                                                                                                                                                                   (UYCO ) UNIV COLUMBIA NEW YORK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9932615-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   screen for modulators and as probes to isolate related genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 40; 60pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          auto:immune
                                                                                                                                                                                                                                                                                                                                           <u>ი</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 Similarity
13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CACTGCACTTACCA 10
                                                                                                                                                            Page 205;
  identified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 6 A; 4 C; 9
                                                                                                                                                                                                                                                                                                                                                                   Grant S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            t with them - for control of apoptosis, e.g. cancer, viral infection, neuro:degeneration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BCNG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98US-0086436.
97US-0997685.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98WO-US27630
                                                                                                                                                                                                              ion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                       213pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      92.9%;
92.9%;
                                                                                                                                                                                                                                                                                                                                                                   Kandel ER,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 13; DB Pred. No. 3.6e 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                Santoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; DB 19;
. 3.6e+02;
                          Compounds, e.g. antagonists
                                                                                                                                                                                                                                                                                                                                                                'n
                                                                                                                                                                                                                                    or
                                                                                                                                                                                                                                    heart cyclic
                                                                                                                                                                                                                                                                                                                                                                Siegelbaum
for modulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                       and
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RESULT 3
ABK44429
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            neurological, renal, pulmonary, hepatic or cardiovascular condition. Such conditions include epilepsy, Alzheimer's Disease, Parkinson's Disease, long QT syndrome, sick sinus syndrome, age-related memory loss, cystic fibrosis, sudden death syndrome or pacemaker rhythm dysfunction. BCNG or BCNG-related protein can also be used to treat sensory disorders, e.g. blindness, loss of vision, loss of smell, numbness and lack of ability to taste. Also treatable are auditory disorders, respiratory disorders due to defects in central nervous system areas that control respiration or defects in the lungs, dyslexia, attention deficit disorder secretions. The proteins are useful targets for screening for drugs that are effective in the control of main and honoral controls.
WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 HGK; NIK; Nck-interacting kinase; infection; inflammat antisense gene therapy; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 27 BP; 6 A; 7 C; 9
                           Dean NM,
                                                      (ISIS-)
                                                                                   29-AUG-2000;
                                                                                                             29-AUG-2000;
                                                                                                                                                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                               modified_base
                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human HPK/GCK-like kinase antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABK44429 standard;
                                                                                                                                          12-FEB-2002
                                                                                                                                                                   US6346416-B1
                                                                                                                                                                                                                                         modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BCNG-related protein activity. Modulation is increased or decapermissivity or ion flow rate. Modulators of BCNG can be used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     effective in the control of pain and hyperalgesia
2002-237091/29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HPK/GCK-like kinase; antiinflammatory; cytostatic; antimicrobial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 CACTGCACTTNCCA 14
                                                      SISI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CACTGCACTTGCCA 8
                          Cowsert LM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                       PHARM INC
                                                                                   2000US-0651011
                                                                                                             2000US-0651011
                                                                                                                                                                                                                                                                                                            /note=
1..20
                                                                                                                                                                                                                                         16..20
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                                                                                                                                                                                                            /*tag=_c
/mod_base=_OTHER
                                                                                                                                                                                                                                                                  /note=
                                                                                                                                                                                                                                                                                /mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  92.9%;
92.9%;
                                                                                                                                                                                                                                                                                                                        "Optionally
                                                                                                                                                                                            "Optionally 2'-methoxyethyl (2'MOE) nucleotides"
                                                                                                                                                                                                                                                   "Phosphorothioate backbone; are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         G; 5 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                      2'-methoxyethyl (2'MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                  all cytidines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ISIS 105328
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          to treat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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New antisense compound, useful for preventing or delaying infection,

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RESULT 4
AAS03611
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The sequence represents PCR primer rend9bl, used to detect RHD positive haplotypes in RHD negative individuals. The primer was used in DNA typing using PCR-sequence length polymorphism (SSP) of the Rhesus genes locus comprising the RHD, SMP1 and RHCE (all undefined) genes and/or the Rhesus box(es), preferably the hybrid Rhesus box, the upstream Rhesus box and/or the downstream Rhesus box. The RHD and RHCE genes are located at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  region of a nucleic acid molecule encoding HPK/GCK (undefined)-like Kinase (HGK) (also known as NIK for Nck-interacting kinase), which specifically hybridises with and inhibits expression of HGK. (I) is useful for inhibiting the expression of HPK/GCK-like kinase in cells or tissues in vitro. (I) is useful prophylactically e.g. to prevent or delay infection, inflammation and tumour formation. (I) is also useful as a diagnostic and research reagent. (I) Is also useful for distinguishing functions of various members of a biological pathway and in antisons of care thereone who present secures a second to the contract of a pathway and in antisons of care thereone who present secures as the contract of the contract
                                                                                                                                                                                                                                                                                                                                                                      New nucleic acid molecular structure, RHD positive haplotypes in D-negative and RHCE genes -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Flegel WA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200132702-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rhesus box; RHD positive; sequence length polymorphism; SSP; RHD; SMP1; RHCE; RHD negative; blood group typing; blood transfusion; antigen C; haemolytic disease of the newborn; chromosome 1 p34.1-p36; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAS03611
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAS03611 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seguence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       oligonucleotide targeted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antisense
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (DRKB-) DRK BLUTSPENDEDIENST BADEN WUERTTEMBERG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-NOV-1999; 99EP-0121686.
31-MAY-2000; 2000EP-0111696.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-OCT-2000; 2000WO-EP10745
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           inflammation or tumour formation, is targeted to nucleic acid molecule encoding HPK/GCK-like kinase (HGK) and hybridises and inhibits HGK
                                                                                                                                                                                                                                                                                          Example 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer rend9b1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2001-291052/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ب
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               vention relates to an antisense compound (I) of 8-50 nucleobases in targeted to a start codon region, coding region or 3'-untranslated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CACTGCACTINCCA 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gene therapy.
                                                                                                                                                                                                                                                                                          Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Wagner FF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 3 A; 7 C; 4 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                          56;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     81.4%;
85.7%;
                                                                                                                                                                                                                                                                                      135pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence represents an antisense d to human HPK/GCK-like kinase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                          English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 detect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 11.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RHD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NO.
                                                                                                                                                                                                                                                                                                                                                                                                   useful for detection of common individuals, comprises RHD, SMP1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        positive haplotypes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ņ
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멍 Qy

Query Match

81 . 48;

Score 11.4;

В 9;

Length 27;

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RESULT 5
AAN81248
ID AAN8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ХX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         chromosome 1 p34.1-p36. Rhesus box flanks the breakpoint region of the RHD deletion in the common RHD negative haplotypes. The primers of the invention are useful for: (1) the specific detection of the common RHD positive haplotypes in D-negative individuals; (2) blood group typing; (3) determining whether a patient can be transfused with RhD negative blood and whether blood is suitable for transfusion to patients who should not be exposed to antigen C; (4) assessing the risk of a RhD negative mother of conceiving or carrying an RhD positive foetus. Antiantibodies are useful for treating pregnant women who are Rhesus D negative, where the foetus is not homozygous for the RHD gene to treat
                                     No. repeat units: 1.

No. repeat units: 1.

Category of hybridisation to human DNA: 1.

Category of hybridisation to human DNA: 1.

The probe is complementary to one or two repeat units of VTNR loci.

The probe is used for genetic identification of a sample of human genomic DNA, eg for paternity testing, diagnosing cancer or genetic diseases, and studying post-bone marrow transplant chimerism. Under high criteria, it yields a locus-specific or mulit-loci, polymorphic hybridisation pattern, and is more specific for a single locus (or small no. of loci) than known probes.
 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic probe; restriction fragment length polymorphism; paternity testing; ss.
                                                                                                                                                                                                                           Claim 7(e); Page 6;
                                                                                                                                                                                                                                                                       New oligonucleotide hybridisation probe
                                                                                                                                                                                                                                                                                                     WPI; 1988-347751/49
                                                                                                                                                                                                                                                                                                                                                                                              17-MAY-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                            26-MAY-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-DEC-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Probe
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                                See also
                                                                                                                                                                                               Total length: 27
                                                                                                                                                                                                                                                       with high specificity for single
                                                                                                                                                                                                                                                                                                                                                               (CITY )
                                                                                                                                                                            Length repeat unit:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                prevent haemolytic disease of the newborn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0-YNH24 to identify intrinsic polymorphism of loci in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 CACTGCACTTNCCA 14
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12; Conserv
                                                                                                                                                                                                                                                                                                                                                                 CITY
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   27
                                N981243-53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 5 A; 7 C; 5
 BP;
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                                                                                                                                                                                                                                                                                                                                                                 OF.
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                                                                                                                                                                                                                                                                                                                                                                                              88US-0194982
                                                                                                                                                                                                                                                                                                                                                                                                                            88EP-0304763
 7 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                               31.
                                                                                                                                                                                                                          9pp; English
12 C;
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85.7%;
                                                                                                                                                                                                                                                                                                                                                                 MED CENTRE
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 2 G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 11.4; D
Pred. No. 3.1e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0,
 6 T; 0 other;
                                                                                                                                                                                                                                                         locus,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.1e+03;
2;
                                                                                                                                                                                                                                                    specific for repeat units useful e.g. in paternity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                                       paternity testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               foetus. Anti-D
Rhesus D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
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RESULT 6
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                                                                                                                                                                           RESULT 7
                                                                                                                                                                                                                                                                   Ouery Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                           The probe is used for genetic identification of a sample of human gene DNA, e.g. for paternity testing, diagnosing cancer or genetic diseases and studying bone marrow transplant chimerism. Under high criteria it yielded locus-specific or multi-loci, polymorphic hybridisation patter and is more specific for a single locus (or small number of loci) than
                       Human; probe; hypervariable number of tandem repeat; VN species identification; virus detection; minisatellite
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New oligonucleotide hybridisation probe specific for repeat units with high specificity for single locus, useful e.g. in paternity
             Epstein-Barr virus;
                                                                                            12-SEP-2001
                                                                                                                                                AAS04026 standard; DNA; 31
                                                                                                                                                                                                                                                                                                                       Sequence 27
                                                                                                                                                                                                                                                                                                                                                   known probes.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 7; page 6; 8pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1988-347751/49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wallace RB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-MAY-1987;
17-MAY-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Probe 0-YNH24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-DEC-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAN82045 standard; DNA;
                                                                  VNTR hybridisation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CITY ) CITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-MAY-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAN82045
                                                                                                                                                                                                                 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14
                                                                                                                                                                                                                                                                    12;
                                                                                                                                                                                                                               1 CACTGCACTTNCCA 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 CACTGCACTTNCCA 14
                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CACTGTACTTCCCA 27
                                                                                                                                                                                                                 CACTGTACTTCCCA 27
                                                                                                                                                                                                                                                                                                                         BP; 7
                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             QF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                           (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      87US-0055224
88US-0194982
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88EP-0304763
                                                                                                                                                                                                                                                                                                                         Α.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human genomic DNA.
                                                                 probe, minisatellite YNH24
                                                                                           entry)
                                                                                                                                                                                                                                                                                                                       12 C;
                                                                                                                                                                                                                                                                                  85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   85.7%;
                                                                                                                                                                                                                                                                                 . 78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          probe 0-YNH24; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                                                                                                         2
G;
                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No. 3.1e+03;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                 Score 11.4;
Pred. No. 3
                                                                                                                                                                                                                                                                                                                       6 T; 0 other;
                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                               .1e+03;
                                                                                                                                                                                                                                                                                             DB
                                                                                                                                                                                                                                                                                             9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                          Length 27;
                                                                                                                                                                                                                                                                                                                                                             hybridisation pattern, number of loci) than
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         human genomic DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                    diseases,
                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             . ;
                                                                                                                                           RESULT 8
AAQ49016
ID AAQ4
                                                                                                                                                                                                                                                                                                                       В
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XXXXXXX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  XX
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            XPX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FTT
                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The sequence represents the nucleotide sequence of variable number of tandem repeats (VWTR) hybridisation probe, minisatellite YNH24. The sequence was used as a probe to demonstrate the method of the invention, which is a device for detecting or characterising a nucleic acid analyte. The device comprises a panel of double stranded oligonucleotide probes immobilised on a solid support. Each probe has a first strand comprising a hypervariable number of tandem repeat sequences and a second strand complementary to the first strand. The probe is selected to extrand the selected to the first strand.
                                                                                                                                                                                                                                                                                                                                                                                                  characterising a nucleic acid analyte without requiring electrophoresis or the direct sequencing of analyte samples or analyte fragments. The device is particularly useful for individual and species identification.
                                                                                                                                                                                                                                                                                                                                                                                                                                        hybridise to at least one allele of the nucleic acid analyte and comprises a fragment of an Epstein-Barr virus genome spanning from nucleotide 7421-8042. The device is useful for detecting or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New device with a panel of double stranded oligonucleotide probes immobblized on a solid support, useful for detecting or characterizing a nucleic acid analyte without requiring electrophoresis or the direct sequencing of analyte
        SBP; specific binding pair members; antibody; RGDP, replicable genetic display package; recombination; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                Sequence 31 BP;
                                                                                                                                                                                                                                                                                                                                                          prior methods. The device is also readily applied outside of a laboratory environment, such as detection of pathogenic virus strains under
                                                               Multimeric
                                                                                          22-APR-1994
                                                                                                                    AAQ49016;
                                                                                                                                             AAQ49016 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                      The device is simple and reliable compared to those previously used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Yeh HR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-APR-1996;
16-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Column 20; 24pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-366471/38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (USSA)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                              9
                                                                                                                                                                                                                                      1 CACTGCACTTNCCA 14
                                                                                                                                                                                                                                                                   Similarity 12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   S
                                                                                                                                                                                                                                                                                                                                                conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEC
                                                                                                                                                                                                                                                                   Conservative
                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   QF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            96US-0015965
97US-0838157
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "Labelled with 4'~6-diamidino-2-phenylindole'
                                                              antibody chain primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ′*taq=
                                                                                                                                                                                                                                                                                                                      Α;
                                                                                                                                                                                                                                                                              81.4%;
85.7%;
                                                                                                                                                                                                                                                                                                                      14 C;
                                                                                                                                                                                                                                                                                                                                                in the
                                                                                                                                             36
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                                                                                                                                                                                                                                                                                                                      ω
                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                                field
                                                                                                                                                                                                                                                                                                                      G; 7 T; 0 other;
                                                                                                                                                                                                                                                                              Score 11.4; DB 22
Pred. No. 3.2e+03;
                                                                                                                                                                                                                                                                                          22;
                                                                                                                                                                                                                                                                   2;
                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                         Length
                          PCR;
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                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                    AAH46243
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                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
                                                                                                                                                 WO200146237-A1
                                                                                                                                                                      Homo
                                                                                                                                                                                          cytostatic; anti-HIV; antiinflammatory;
                                                                                                                                                                                                                                             Human neuronal thread
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                               The primers (AAQ48987-Q49045) are used in the amplification of Kappa and lambda-chain genes of various antibodies. These genes are then recombined into the same replicon, resulting in very diverse librar: of antibody chains, e.g. from unimmunised donors. It is also useful for chain shuffling, mutagenesis, humanising and CDR imprinting.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 58; 81pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Prodn. of specific binding pair members, e.g. antibody chains by display on surface of replicable genetic display packages % \left( 1\right) =\left( 1\right) ^{2}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Griffiths AD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
  Human neuronal thread protein 17 and
                                                                                     22-DEC-1999;
                                                                                                                                                                                                               malignant
                                                                                                                                                                                                                                                                                                           AAH46243 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-MAR-1992;
15-MAY-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30-SEP-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              W09319172-A.
                                                                                                       18-DEC-2000;
                                                                                                                                                                                                                                                                  25-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MEDI-) MEDICAL RES. COUNCIL.
                                                                (BIOW-)
                                                                                                                                                                                                                        Human neuronal thread protein 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CAMB-) CAMBRIDGE ANTIBODY TECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                           ۲,
                      2001-441667/47.
                                                                                                                                                                      sapiens
                                                                                                                                                                                                 nant tumour; cancer; blood immunodeficiency virus; in
                                                                                                                                                                                                                                                                                                                                                               9
                                                                                                                                                                                                                                                                                                                                                                                 1 CACTGCACTTNCCA
                                           Xie
                                                                BIOWINDOW GENE DEV INC
                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                              CAGTGCACTTGCCA
                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 10 A; 13 C; 7 G; 6 T;
                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                         2000WO-CN00585
                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     92GB-0006318
92WO-GB00883
                                                                                     99CN-0125691
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Johnson KS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     93WO-GB00605
                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                81.4%;
85.7%;
                                                                                                                                                                                                                                            protein 17
                                                                                                                                                                                                                                                                                                           41
                                                                                                                                                                                                                                                                                                                                                              22
                                                                                                                                                                                                                                                                                                           ВP
                                                                                                                                                                                                                                                                                                                                                                                                      0,:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Smith AJH,
                                                                                                                                                                                                                                                                                                                                                                                                                         Score 11.4;
                                                                                                                                                                                                  immune disorder;
                                                               SHANGHAI
                                                                                                                                                                                              mmune disorder; inflammatory condition;
                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                           probe,
                                                                                                                                                                                                                        recombinant production;
                                                                                                                                                                                                                                                                                                                                                                                                                 No.
encoded polynucleotide, applicable
                                                                                                                                                                                                                                                                                                                                                                                                                                               0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                3.3e+03;
                                                                                                                                                                                       immunomodulator; probe;
                                                                                                                                                                                                                                            SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Waterhouse
                                                                                                                                                                                                                                                                                                                                                                                                                         DB 14;
                                                                                                                                                                                                                                             NO:8
                                                                                                                                                                                                                                                                                                                                                                                                                         Length
                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                     0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       libraries
                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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AAA66165
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      ₹ ₹ ₹
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  production of the protein. The present invention additionally discloses an antagonist of neuronal thread protein 17 for therapeutic use, and an antibody which specifically binds to neuronal thread protein 17. Neuronal thread protein 17. and nucleotides which encode it may be used for treating a variety of diseases, such as malignant tumours, blood diseases, HIV (human immunodeficiency virus) infection, immune disorders and inflammatory conditions. The protein may also be used to screen for modulators of its activity or for peptide fingerprinting identification. The polynucleotide can be used as a primer for nucleic acid amplification reactions or as a probe for hybridisation reactions, or in producing gene chips or microarrays. Sequences AAH46243-AAH46244 represent human reported thread protein 17 probes used in an exemplification of the
(Canine familiaris) genome comprising the genome location of a marker selected from AAA66139 to AAA66942. The radiation hybrid map is useful for identifying and localising dog genes, since it covers approximatel 80 % of the dog genome and provides a dense map integrating different
                                                                                                                           New radiation hybrid map of the dog, Canine familiaris, genome, useful for e.g. identifying genes implicated in phenotypic and behavioral traits or in genetic diseases and for studying dog pedigrees \,\cdot\,
                                                                                             Claim 1; Page 54; 87pp; English
                                                                                                                                                                                                                           Galibert F,
                                                                                                                                                                                                                                                                                                                                                                                                                       Canis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dog; genome; genomic marker; radiation hybrid chromosome location; gene marker; polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dog genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 41 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               in diagnosis and treatment of cancer, immunological diseases and phlogosis
                                                               The present invention describes a radiation hybrid
                                                                                                                                                                                                                                                                                          13-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                        25-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                       WO200029615-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                   phenotype;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA66165;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA66165 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention
                                                                                                                                                                                                                                                           (CNRS ) CNRS
                                                                                                                                                                                                                                                                                                                          15-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                     familiaris.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 CACTGCACTTNCCA 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CACTGCAATTTCCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                      behaviour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  marker oligonucleotide sequence SEQ ID NO:27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                           Andre C;
                                                                                                                                                                                                                                                           CENT NAT RECH SCI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                relates to human neuronal thread protein 17 (AAB98894), encoding it (AAH46238), and a method for the recombinant
                                                                                                                                                                                                                                                                                          98US-0108193
                                                                                                                                                                                                                                                                                                                          99WO-IB01907
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Α.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          81.4%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      pedigree; ss
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Pred. No. 3.3e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Τ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hemopathy, HIV infection,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .3e+03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    map; identification;
microsatellite marke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
                                                               map
 covers approximately 
regrating different
                                                               of the dog
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            41;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     marker;
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RESULT 11
ABI94202/c
ID ABI942
XX ABI942
XX ABI942
XX Human:
Captur
XX Human:
KW ligase
KW cancer
KW cancer
KW cancer
XX Synthe
PN W02001
XX Gorde
PN W02001
XX Haran
YX Haran
YX Haran
YX Haran
YX CORR
YX Human:
KW Cancer
XX Synthe
YX Concer
XX Synthe
YX Gorde
YX Haran
YX Haran
YX Haran
YX CORR
YX CORR
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YX Go
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Best Local
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                       e.g. Salmoneila, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents e.g. Cryptococcus neoformans, Candida albicans and Aspergillus funigatutus, viruses e.g. Troell lymphocytotrophis cirus, Epstein-Barr virus and polio virus, and parasitic infectious agents selected from Onchoverva volvulus, Entameba histolytica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obesity defects. Detecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, passon and papillomavirus types 16 and 18 and liver cancers. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes a method (M1) for designing capture oligonucleotide probes (I) for use on a support to which complementary oligonucleotide probes (II) will hybridise with little mismatch, where
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; K-ras; PCR primer; probe; capture probe; mutation detection; ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious diseas infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic; environmental monitoring; food industry; feed industry; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 types (i.e. Type I and Type II) of markers. The map and the do markers (or complementary sequences) are especially useful to genes responsible for phenotypic and behavioural traits in dog identify morbid genes, to analyse diseases and identify implicant such diseases and their alleles, and to study dog pedigrees may also be useful for isolating corresponding human gene seque.g. genes involved in genetic diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch \boldsymbol{\cdot}
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                                                                                                                                                                                                                                                                                                                                                                                   (1) have melting temperatures within a narrow range. The method is useful
for detecting infectious diseases caused by bacterial infectious agents
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   dene,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29; 300pp; English.
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   for environmental monitoring,
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Pred. No.
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to identify
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Best Local :
                                                             e.g. Salmoneila, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents e.g. Cryptococcus neoformans, Candida albicans and Aspergillus funiquatus, viruses e.g. T-cell lymphocytotrophis cirus, Epstein-Barr virus and polio virus, and parasitic infectious agents selected from Onchoverva volvulus, Entamoeba histolytica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obesity defects. Detecting cancer involving oncogenes, tumour suppressor genes, or genes
involved in DNA amplification, replication, recombination or repair, cancer is specifically associated with a gene selected from BRCA1 gene, human papillomavirus types 16 and 18 and liver cancers. The method is also used for environmental modification.
                                                                                                                                                                                                                                              oligonucleotide probes (I) for use on a support to which complementary oligonucleotide probes (II) will hybridise with little mismatch, where
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; fore environmental monitoring; food industry; feed industry; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and feed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the particular sites and identifying if ligation of the oligonucleotide probe sets occurred and correlating (using a computer) identified ligation to a presence or absence of the target nucleotide sequences. ABI82074 to ABI97366 represent oligonucleotide sequences used in the exemplification
                                                                                                                                                                                                                                                                                                                                                     Designing capture oligonucleotide probes complementary oligonucleotides hybridize

    have melting temperatures within a narrow range. The method is useful
for detecting infectious diseases caused by bacterial infectious agents

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Capture oligonucleptide Zip ID#3885 oligo #9.
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                                                                                                                                                                                                                                                                                  present invention describes a method (M1) for designing capture
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          K-ras; PCR primer; probe; capture probe; mutation detection; detection reaction; LDR; p53; BRCA1; BRCA2; infectious disea
                                                                                                                                                                                                                                                                                                                     5; Fig 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Zirvi M,
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91.7%;
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                                    selected from BRCAl gene,
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                                                                                                                                                 for detecting infectious diseases caused by bacterial infectious agents e.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents e.g. Cryptococcus neoformans, Candida abbicans and Aspergillus fumigautus, viruses e.g. T-cell lymphocytotrophis cirus, Epstein Barr virus and polio virus, and parasitic infectious agents selected from Onboverva volvulus, Entamoeba histolytica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obesity defects.
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                                                                                                                                                                                                                                                                                                                                                                                                         oligonucleotide probes (I) for use on a support to which complementary oligonucleotide probes (II) will hybridise with little mismatch, where (I) have melting temperatures within a narrow range. The method is useful
                                                                                            Detecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Designing capture oligonucleotide probes complementary oligonucleotides hybridize
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Barany F, Zirvi M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-APR-2000; 2000US-197271P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Capture oligonucleotide Zip ID#1289 oligo
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is specifically associated with a gene selected from BRCAl gene, ne, human papillomavirus types 16 and 18 and liver cancers. The is also used for environmental monitoring, forensics and the food
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                                                                            for detecting infectious diseases caused by bacterial infectious agents e.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, funga infectious agents e.g. Cryptococcus neoformans. Candida abbicans and Aspergillus fumigautus, viruses e.g. T-cell lymphocytotrophis cirus, Epstein Barr virus and polio virus, and parasitic infectious agents selected from Onchoverva volvulus, Entamoeba histolytica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency. Turner Syndrome and obesity defects.
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                                                                                                                                                                                                                    oligonucleotide probes (I) for use on a support to which complementary oligonucleotide probes (II) will hybridise with little mismatch, where (I) have melting temperatures within a narrow range. The method is useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; K-ras; PCR primer; probe; capture probe; mutation detect ligase detection reaction; LDR; p53; BRCA1; infectious d infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; fore environmental monitoring; food industry; feed industry; ss.
          Detecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, p53 gene, human papillomavirus types 16 and 18 and liver cancers. The
                                                                                                                                                                                                                                                                                                                                            Designing capture oligonucleotide probes for use on a support complementary oligonucleotides hybridize with little mismatch
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                                                                                                                                                                                                                                                                        present invention describes a method (M1) for designing capture
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91.7%;
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                                                                              (I) have melting temperatures within a narrow range. The method is useful for detecting infectious diseases caused by bacterial infectious agents e.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents e.g. Cryptococcus neoformans, Candida albicans and Aspergillus fumigautus, viruses e.g. T-cell lymphocytotrophis cirus, Epstein Barr virus and polio virus, and parasitic infectious agents selected from Onchoverva volvulus, Entamoeba histolytica and Bracunculus medinesis. The method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obesity defects. Detecting cancer involving oncogenes, tumour suppressor genes, or genes
                                 involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, p53 gene, human papillomavirus types 16 and 18 and 10....
                                                                                                                                                                                                                                                                  The present invention describes a method (M1) for designing capture oligonuclectide probes (I) for use on a support to which complementary oligonuclectide probes (II) will hybridise with little mismatch, where
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Barany F, Zirvi M,
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                               human papillomavirus types 16 and 18 and
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91.7%;
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                             liver cancers.
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                                                                                                                                            and feed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the particular sites and identifying if lightion of the oligonucleotide probe sets occurred and correlating (using a computer) identified lightion to a presence or absence of the target nucleotide sequences. ABI82074 to ABI97546 represent oligonucleotide sequences used in the exemplification
                                                                                                      Sequence
                                                    Local Similarity
                                                                                                                                       the
1 CACTGCACTTNC 12
                                                                                                                                    present invention.
                                                                                                      24
                                   Conservative
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                                                   78.6%;
91.7%;
                                   0; Mismatches
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Pred. No. 5
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Search completed: June 21, 2003, 22:26:25 Job time: 79.7812 secs

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Minimum DB seq length: 0 Maximum DB seq length: 50
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Perfect score:
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   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
 hits satisfying chosen parameters:
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14
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/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
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/cgn2_6/ptodata/1/ina/backfiles1.seq:*
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US-09-261-011k-29
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US-09-260-260a-83
US-08-350-260a-120
US-08-350-260a-120
US-09-050-783-37
US-08-350-706a-2
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Query Match 92.9%; Score 13; Best Local Similarity 92.9%; Pred. No.	RESULT 1  US-08-713-557B-23/c  Sequence 23, Application US/08713557B  Patent No. 5912168  PALECANT: Watson, James D. APPLICANT: Watson, James D. APPLICANT: Watson, CD95 REGULATORY GENE TITLE OF INVENTION: CD95 REGULATORY GENE TITLE OF INVENTION: AND TRANSCRIPTION FA  PARTICANT: Watson, CD95 REGULATORY GENE TITLE OF INVENTION: AND TRANSCRIPTION FA  COMPUTER OF SEQUENCES: 38  CORRESPONDENCE ADDRESS: ADDRESSEE: LAW Offices of Ann W. Speck STATE: WA  CITY: Seattle COUNTRY: USA ZIP: 98121  COMPUTER READABLE FORM: MEDIUM TYPE: Diskette COMPUTER READABLE FORM: MEDIUM TYPE: Diskette COMPUTER: IBM Compatible OPERATION STEEN APPLICATION NUMBER: US/08/713,557B FILING DATE: 30-AUG-1996 CLASSIFICATION NUMBER: US/08/713,557B FILING DATE: 30-AUG-1996 CLASSIFICATION NUMBER: 11000.1004 TELEPHONE: SPECKMAN, AND W REGISTRATION NUMBER: 11000.1004 TELEPHONE: 206-269-0565 TELEPHONE: 206-269-0565 TELEPHONE: 206-269-0565 TELEPHONE: 206-269-0565 TELEPHONE: 206-269-0565 TELEPHONE: 206-269-0565 TELEPHONE: Shage pairs TYPE: nucleic acid STRANDEDNESS: Single TOPOLOGY: linear US-08-713-557B-23	28 10.4 74.3 41 1 US-08-307-619-61 29 10.4 74.3 41 2 US-08-350-260A-10 30 10.4 74.3 41 2 US-08-350-260A-20 31 10.4 74.3 41 3 US-09-050-783-61 31 10.4 74.3 44 3 US-09-050-783-61 32 10.71.4 20 1 US-08-564-002-1 34 10 71.4 20 1 US-09-180-437-11 36 10 71.4 20 4 US-09-180-437-231 37 10 71.4 20 4 US-09-180-437-231 38 10 71.4 20 4 US-09-180-437-231 39 10 71.4 21 US-08-136-741-7 39 10 71.4 21 US-08-136-741-7 39 10 71.4 22 4 US-09-460-145-6 40 10 71.4 23 US-09-000-094-1 50 10 71.4 30 4 US-09-000-094-1 60 10 71.4 30 4 US-09-000-094-1 61 10 71.4 31 US-08-169-715-47 62 10 71.4 31 US-08-169-715-47 63 10 71.4 31 US-08-169-718-35 64 10 71.4 31 US-08-169-718-35
DB 2; Length 25;	GENE SEQUENCES ON FACTORS Speckman 4185 4185	619-61  260A-107  Sequence 61, Appl 260A-208  Sequence 208, App 260A-583  Sequence 1, Appl 347-11  347-11  347-11  347-10  Sequence 1, Appl 384A-70  Sequence 11, Appl 384A-70  Sequence 21, Appl 437-231  Sequence 70, Appl 437-231  Sequence 177, App 741-7  Sequence 37, Appl 145-6  Sequence 6, Appli 145-37  Sequence 37, Appl 145-47  Sequence 39, Appl 715-47  Sequence 39, Appl 715-47  Sequence 39, Appl 715-47  Sequence 39, Appl 716-35  Sequence 37, Appl 716-35  Sequence 39, Appl 716-37  Sequence 39, Appl 716-38  Sequence 37, Appl 716-39  Sequence 37, Appl 716-47  Sequence 39, Appl 716-47  Sequence 39, Appl 716-47  Sequence 37, Appl 716-35  Sequence 37, Appl 716-47  Sequence 37, Appl 716-48  Sequence 37, Appl 716-48  Sequence 37, Appl

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 7, Application Patent No. 6238866
. GENERAL INFORMATION: Yeh,
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SEQ ID NO 29
LENGTH: 20
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF HPK/GCK-LIKE KINASE EXPRESSION
FILE REFERENCE: RTS-0168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/651,011A CURRENT FILING DATE: 2000-08-29
                                                                                                                                                            TELEFAX: (410) 436-2534 INFORMATION FOR SEQ ID NO: 7:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
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nes 12; Conserv
                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 31 base pairs
                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 436-96
TELECOMMUNICATION INFORMATION:
TELEPHONE: (410) 436-1158
                                                                                                                                                                                                                                                                                                   CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Office of the Chief Counsel
STREET: U.S. Army SBCCOM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Yeh, Homer R., Wick, Charles H.
TITLE OF INVENTION: NOVEL DETECTOR FOR NUCLEIC ACID
TYPING AND METHODS OF USING THE SAME.
HYPOTHETICAL:
                                     MOLECULE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 CACTGCACTTNCCA 14
                                                                           TYPE: nucleic acid
STRANDEDNESS: not relevant
                                                            TOPOLOGY: not
                                                                                                                                                                                                                                                            NAME: Biffoni, U. John REGISTRATION NUMBER: 39908
                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/09/246,277A FILING DATE: 08-Feb-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: Maryland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: APG(EA)
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                                   TYPE: DNA (genomic)
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85.7%;
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RESULT 4
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CHROMOSOME/SEGMENT: YNH24 Minisatellite
SEQUENCE DESCRIPTION: SEQ ID NO: 7:
US-09-246-277A-7
Query Match
Best Local Similarity
Matches 12; Conserv
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                                                                                                                                                                                     REGISTRATION NUMBER: 36,107
REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                APPLICATION NUMBER: PCT/G
FILING DATE: 15-MAY-1992
ATTORNEY/AGENT INFORMATION:
NAME: David W. Clough
                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: GB 9: FILING DATE: 24-MAR-1992 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
CLASSIFICATION: G01N
PRIOR APPLICATION DATA:
                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 24-MAR-1993 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
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                                                                                          STRANDEDNESS:
                                                                                                                                                                             TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE:
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                                                                                                                           LENGTH:
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12; Conservative
                                                                                                           nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Johnson,
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Smith, Andrew JH
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PatentIn Release #1.0, Version #1.25 (EPO)
                                                                               linear
                                                                                                                                                                         312-474-6300
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              81.4%;
85.7%;
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85.7%;
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                                                                                                                                                          37:
              Score 11.4; DB 1; Pred. No. 4.3e+02;
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Pred. No. 4.2e+02;
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                             DB 1;
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                            Length 36;
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Matches

12;

Conservative

0;

Pred. No. 4.3e+02; 0; Mismatches 2;

Indels

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Gaps

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85.78;

Best Local Similarity

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RESULT 5
US-08-350-260A-83
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                                   US-08-350-260A-83
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Query Match
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                                                                                                                                 NFORMATION FOR SEQ ID NO:
                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                             FILING DATE: 24-MAR-1993 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: GB 9
FILING DATE: 24-MAR-1992
PRIOR APPLICATION DATA:
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                                                                                                                                           TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-474-6300
                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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APPLICATION NUMBER:
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STREET:
CITY: C
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OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 01 FILING DATE: 31-MAR-1994
                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: PCT/O
                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: PCT/
FILING DATE: 10-JUL-1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
ZIP: 60606-6402
                                                                                                                                                                                               NAME: Clough, David W
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE:
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                                                               STRANDEDNESS:
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                                                                                                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
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                                                                                               LENGTH:
                                                           H: 36 base pairs nucleic acid NDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chicago
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Marshall, O'Toole, Gerstein, Murray & Borun
6300 Sears Tower, 233 South Wacker Drive
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VENTION: Methods for producing members of specific
VENTION: binding pairs
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                 linear
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 81.4%;
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 Score 11.4;
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                                                       NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                 NAME: CLOUGH, DAVID W
REGISTRATION NUMBER: 36,107
REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9110549.4
FILING DATE: 15-MAY-1991
PRIOR APPLICATION DATA:
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CURRENT APPLICATION DATA:
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APPLICATION NUMBER:
FILING DATE: 24-MAR-
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                                     SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                          APPLICATION NUMBER: FILING DATE: 31-MA
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OPERATING SYSTEM: PC-DOS/MS-DOS
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STREET: Marshall, O'Toole, Gerstein, Murray & Borun
STREET: 6300 Sears Tower, 233 South Wacker Drive
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                                                                       TELEPHONE:
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nucleic acid
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VENTION: Methods for producing members of specific
VENTION: binding pairs
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                                                                       312-474-6300
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31-MAR-1994
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                                                        Matches
                                                                                   Query Match
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Best Local Similarity
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SEQUENCE CHARACTERISTICS:
LENGTH: 36 base
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                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                            FILING DATE: 15-MAY-1992
ATTORNEY/AGENT INFORMATION:
NAME: David W. Clough
REGISTRATION NUMBER: 36,
                                                                                                                                                                                                                                                                                                                     FILING DATE: 24-MAR-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/
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FILING DATE: 24-MAR-
PRIOR APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 6 CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 16-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
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                                                                                                                            STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                                            TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
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                                                        12;
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                           1 CACTGCACTTNCCA 14
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                                                                                                                                                           nucleic acid
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6140471
CAGTGCACTTGCCA 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USA
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                                                      Conservative
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Griffiths,
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                                                                                                                                          single
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                                                                  81.4%;
85.7%;
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85.7%;
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36,107
APR: 28111/32238
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                                                                    Score 11.4; DB 3; Pred. No. 4.3e+02;
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Pred. No. 4
                                                      Mismatches
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                                                                                  DB 3;
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                                                     Indels
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US-08-362-706A-2
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                                                                                                                                                                                                                                                                                                           RESULT 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                          Sequence 2, Application US/09549808 Patent No. 6338944
                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                           GENERAL INFORMATION:
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MOLECULE TYPE: CDN
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INFORMATION FOR SEQ ID NO
          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Heller Ehrman White & McAuliffe
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LENGTH: 18 base pairs
TYPE: nucleic acid
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REFERENCE/DOCKET NUMBER: 13
TELECOMMUNICATION INFORMATION:
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FILING DATE: 23-DEC-1994
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Schmonsees, William
                                                                                                                                                                                                           APPLICANT: Borden, APPLICANT: Heller,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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APPLICANT: Heller,
TITLE OF INVENTION:
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                                                                                                                                                                            NUMBER OF SEQUENCES:
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                                                          COUNTRY: USA
ZIP: 94301-1900
                                                                                                STATE:
                                                                                                              ADDRESSEE: Heller comments
STREET: 525 University Avenue
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ZIP: 94301-1900
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 OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Palo Alto
                                                                                                                                                                                                                                                                                                                                                        CACTGCCCTTACC 15
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(650)324-0638
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PC-DOS/MS-DOS
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Human Stromelysin-1 Promoter
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US-09-549-808-2
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Best Local Similarity
Matches 11; Conserv
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                NFORMATION FOR SEQ ID NO:
                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: U5/09/021,701
                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
                                            TELECOMMUNICATION INFORMATION: TELEPHONE: 650-236-2386
                                                                                                                       ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
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SEQUENCE CHARACTERISTICS:
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                                                                         NAME: Choi, Wendy A. REGISTRATION NUMBER: 36 REFERENCE/DOCKET NUMBER:
                                                                                                                                    APPLICATION NUMBER: FILING DATE: 10-FEE CLASSIFICATION:
                               TELEFAX:
                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                             STREET: 3000 Har CITY: Palo Alto
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3000 Hanover Street
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                                                                                                                                                                                                                                                                                                 USA
                             650-852-8063
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Wolber, Paul K.
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                                                                                                                                                                                                                                                                                                                                                                                                          probe sequences
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                                                                                        36,697
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Pred. No. 1.
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; ANTI-SENSE: NO
US-09-021-701-469
                                                                               US-09-021-701-470
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                                                    Query Match
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GENERAL I
                            Best Local
Matches
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Best Local
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INFORMATION FOR SEQ ID NO:
                                                                                          MOLECULE TYPE: C
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Choi, Wendy A.
REGISTRATION NUMBER: 36,697
                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION TELEPHONE: 650-236-2386
                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
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                                                                                                                                  TOPOLOGY:
                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Palo Alto
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                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                      IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                  linear
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84.6%;
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1165
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                                       Score 10.4; DB 4; Pred. No. 1.5e+03;
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Pred. No. 1
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                           Mismatches
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                                                    DB 4;
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                          Indels
                          0;
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Best Local Similarity
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                                                                                                                                                                                    GENERAL INFORMATION:
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ENFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE:
HYPOTHETICAL: N
ANTI-SENSE: NO
                                                                APPLICANT: Delenstarr, Glenda C.
APPLICANT: Webb, Peter G.
APPLICANT: Kincaid, Robert H.
TITLE OF INVENTION: Methods for evaluating oligonucleotide
TITLE OF INVENTION: probe sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-236-2386
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NAME: Choi, Wendy A.
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APPLICATION NUMBER: US,
FILING DATE: 10-FEB-19
                                  CORRESPONDENCE ADDRESS:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
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Wolber, Paul K.
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                                                                                                                                                  Wolber, Paul
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                                                probe sequences
1165
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84.6%;
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Manager, Legal Department, Hewlett-Packard Company M/S 20 er Street
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Pred. No. 1.
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RESULT 14
US-09-021-701-473/c
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                                                                                                                                                                                                                         STREET: 3000
CITY: Palo Alto
STATE: CA
                                                                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 10-FEB-19
                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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HYPOTHETICAL: I
ANTI-SENSE: NO
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                                            ATTORNEY/AGENT INFORMATION:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
              NAME: Choi, Wendy A. REGISTRATION NUMBER: 36,697
                                                             CLASSIFICATION:
                                                                                                                           SOFTWARE:
                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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REFERENCE/DOCKET NUMBER:
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                                                                                                                           PatentIn Release #1.0, Version #1.30
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                                                                                                                                                                                                                                                                                                                                                                                              Delenstarr, Glenda C
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84.68;
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1165
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                                                                                            US/09/021,701
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10971464-1
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; MOLECULE TYPE: (
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
US-09-021-701-474
Query Match
Best Local Similarity
Watches 11; Conserve
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US-09-021-701-474/c
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APPLICATION NUMBER: US/09/021,701
FILING DATE: 10-FEB-1998
CLASSIFICATION: PEB-1998
ATTORNEY/AGENT INFORMATION:
NAME: Choi, Wendy A.
REGISTRATION NUMBER: 36,697
REGISTRATION NUMBER: 10971464-1
TELECOMMUNICATION INFORMATION:
TELEPAX: 650-236-2386
TELEPAX: 650-238-2865
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Best Local Similarity
Matches 11; Conserv
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GENERAL INFORMATION:
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[INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: 650-236-2386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Kincaid, Robert H.
TITLE OF INVENTION: Methods for evaluating oligonucleotide
TITLE OF INVENTION: probe sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 20 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: 650-852-8063
                                                                                                                                                TOPOLOGY:
                                                                                                                                                           STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Records Manager, Legal Department, Hewlett-Packard Company M/S 20 STREET: 3000 Hanover Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 94304
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84.6%;
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                   0;
                Score 10.4; DB 4;
Pred. No. 1.5e+03;
0; Mismatches 2;
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                0;
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Search completed: June 22, 2003, 00:01:34 Job time: 18.625 secs

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Result
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: /cgn2_6/ptodata/2/pubpna/U
2: /cgn2_6/ptodata/2/pubpna/U
3: /cgn2_6/ptodata/2/pubpna/U
4: /cgn2_6/ptodata/2/pubpna/U
5: /cgn2_6/ptodata/2/pubpna/U
5: /cgn2_6/ptodata/2/pubpna/U
6: /cgn2_6/ptodata/2/pubpna/U
8: /cgn2_6/ptodata/2/pubpna/U
9: /cgn2_6/ptodata/2/pubpna/U
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /cgn2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*
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/cgn2_6/ptodata/2/pubpna/PCTUS_PUBCOMB.seq:*
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0 US-09-964-666-12

0 US-09-964-412-12

0 US-10-105-004-97

US-10-105-004-97

US-10-215-112-2203

US-10-098-263B-65254

US-09-784-674-479

US-09-784-674-471

US-09-784-674-473

US-09-784-674-475

US-09-784-674-475

US-09-784-674-475

US-09-784-674-475

US-09-784-674-475

US-09-784-674-475

US-09-784-674-475

US-09-784-674-475

US-09-887-145-27

US-09-887-145-27

US-10-115-112-4856

US-10-215-112-4856
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (without alignments)
496.907 Million cell updates/sec
                                                                                           Sequence 12, Appl Sequence 12, Appl Sequence 12, Appl Sequence 203, App Sequence 409, App Sequence 470, App Sequence 471, App Sequence 472, App Sequence 471, App Sequence 473, App Sequence 473, App Sequence 474, App Sequence 475, App Sequence 475, App Sequence 475, App Sequence 475, App Sequence 476, App Sequence 476, App Sequence 476, App Sequence 476, App Sequence 477, App Sequence 476, App Sequence 476
   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Description
          30929,
                                   App1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 1
US-09-964-667-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: DC
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	74.3	74.3	74.3	74.3	74.3	74.3	74.3	74.3	74.3	74.3	74.3	74.3	74.3	74.3	74.3	74.3	74.3		74.3	74.3		74.3	74.3	74.3	74.3	74.3
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	-09-902-	US-09-903-786-275	US-09-903-749A-275	-903-2	US-09-906-700-275	US-09-906-646-275	-786-2	US-09-909-204-275	US-09-904-859-275	-820-2	US-09-907-942-275	US-09-907-613-275	US-09-906-838-275	US-09-906-742-275	US-09-904-011-275	US-09-907-841-275	US-09-907-824-275	US-09-902-853-275	US-09-905-291A-275	US-09-988-899-65	US-10-113-794A-3	US-10-121-857-80	2	US-10-098-263B-94806	US-10-098-263B-53671	US-10-098-263B-48658
	275,	Sequence 275, App	275,	275,		275,	Sequence 275, App	275,	275,	275,	Sequence 275, App	275,	Sequence 275, App	275,	275,	Sequence 275, App	275,	275,	Sequence 275, App	e 65,	3, 7			Sequence 94806, A	Sequence 53671, A	Sequence 48658, A

ALIGNMENTS

```
Sequence 12, Application US/09964667
Publication No. US20030033621A1
GENEAL INFORMATION:
APPLICANT: de la Monte, Suzanne
                                                                                                                          INFORMATION FOR SEQ ID NO: 12: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/964,667
TOPOLOGY: linear MOLECULE TYPE: cDNA SEQUENCE DESCRIPTION: SEQ ID NO: 12:
                                                                                                                                                                                                       NAME: Esmond, Robert W.
REGISTRATION NUMBER: 32,8
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Transgenic Animals and Cel
Screening Drugs Effective
of Alzheimer's Disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 14 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                         LENGTH: 14 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                TELEPHONE: 202-371-2600
TELEFAX: 202-371-2540
                                                                                                                                                                                                                                                                                                              FILING DATE: 28-Sep-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Sterne, Kessler, Goldstein STREET: 1100 New York Ave., Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 20005-3934
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Wands, Jack R.
                                                                                                                                                                                                                                                      32,893
                                                                                                                                                                                                                                 0609.4370000
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Treatment
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RESULT 3
US-09-964-412-12
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US-09-964-666-12
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                                                                                                                                                                                                                                                                                                                 Best Loc
Matches
                                                                           Sequence 12, Application US/0964412
Patent No. US20020129391A1
GENERAL INFORMATION:
APPLICANT: de la Monte, Suzani
                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 12, Application US/09964666 Patent No. US20020104108A1 GENERAL INFORMATION:
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Best Local
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                                                                                                                                                                                                                                                                                                                 Local Similarity 100 hes 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Esmond, Robert W.
RECISTRATION NUMBER: 32,893
REFERENCE/DOCKET NUMBER: 0609.4370000
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
TELEFAX: 202-371-2540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox, P.L.L.C.
STREET: 1100 New York Ave., Suite 600
APPLICANT: de la Monte, Suzanne
Wands, Jack R.
TITLE OF INVENTION: Transgenic Animals and Cell Lines for
Screening Drugs Effective for the Treatment or Prevention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: BPPLICATION NUMBER: US/09/964,666
                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Transgenic Animals and Cell Lines for
Screening Drugs Effective for the Treatment or Prevention
of Alzheimer's Disease
                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE DESCRIPTION: SEQ ID NO: 12:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 28-Sep-2001 CLASSIFICATION: <Unknown>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Washington
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Wands, Jack R.
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Pred. No. 3.3e+
                                                                                                                                                                                                                                                                                                                 Mismatches
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US-10-105-004-97/c
; Sequence 97, Application US/10105004
; Publication No. US20030105002A1
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TELEFAX: 202-371-2540
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                COUNTRY: USA
ZIP: 02109-2170
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/10/105,004
FILING DATE: 22-Mar-2002
PRIOR APPLICATION UNMERE: US/08/754,477
FILING DATE: 22-NOV-1996
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PATCHCIT Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: FOLEY, HOAG & ELIOT LLP
STREET: One Post Office Square
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: <Unknown> ATTORNEY/AGENT INFORMATION:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler,
STREET: 1100 New York Ave.,
                  ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: RIEG COMPOSITIONS AND THERAPEUTIC
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                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14;
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                                                                                                                                                                                                                                                                                                             CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 14 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Esmond, Robert W.
REGISTRATION NUMBER: 32,893
REFERENCE/DOCKET NUMBER: 0609.4370000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 28-Sep-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                          AND DIAGNOSTIC USES THEREFOR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 13; DB 10;
Pred. No. 3.3e+02;
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Length 14; Indels

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                                                                                                                                                                                                                                                                                                     US-10-098-263B-65254
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 14936
SOFTWARE: FRATSEQ for Windows Version 4.0
SEQ ID NO 2203
LENGTH: 25
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                            Sequence 65254, Application US/10098263B Publication No. US20030104410A1 GENERAL INFORMATION:
                                     SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1 SEQ ID NO 65254 LENGTH: 25'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                APPLICANT: Mittman, Michael
TITLE OF INVENTION: Human Microarray
FILE REFERENCE; 3118.1
CURRENT APPLICATION NUMBER: US/10/098,263B
CURRENT FILING DATE: 2003-01-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Michael Mittmann
TITLE OF INVENTION: Method of Genetic Analysis of Probes:
TITLE OF INVENTION: Test3
                                                                                        PRIOR APPLICATION NUMBER: 60/276,759
PRIOR FILING DATE: 2001-03-16
NUMBER OF SEQ ID NOS: 131066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/215,112 CURRENT FILING DATE: 2002-08-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: 3119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Synthetic Oligonucleotide
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                       TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
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4 CACGGCACTITCCA 17
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TYPE: nucleic acid
STRANDEDNESS: single
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85.7%;
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85.7%;
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Pred. No. 2
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RESULT 7
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Best Local Similarity 85.
Matches 12; Conservative
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                                                                   Matches
                                                                                    Query Match
Best Local :
                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 469: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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TELEPHONE: 650-236-2386
TELEFAX: 650-852-8063
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 94304
COMPUTER READABLE FORM:
                                                                                                                                                       ANTI-SENSE: NO SEQUENCE DESCRIPTION: SEQ ID NO: 469:
                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Shannon,
                                                                                                                                                                                          HYPOTHETICAL: NO
20 ACTGCATTTACCA 8
                                                                   11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8 CACTGCACTTACAA 21
                                2 ACTGCACTTNCCA 14
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                                                                                      Similarity
                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Cho1, Wendy A. REGISTRATION NUMBER: 36,697
                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 09/021,701 FILING DATE: 10-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/09/784,674
FILING DATE: 15-Feb-2001
CLASSIFICATION: No. US20030054346A1 available
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 3000 Hanover Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                              LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kincaid, Robert H. OF INVENTION: Methods for evaluating oligonucleotide {\sf OF}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/09784674
                                                                   Conservative
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Delenstarr, Glenda C
Webb, Peter G.
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Company M/S 20BO
                                                                                    74.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,:
                                                                 Score 10.4; DB Pred. No. 1e+04; 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Version #1.30
                                                                                                  9;
                                                                                                 Length 20;
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                                                                   Indels
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US-09-784-674-470/c

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RESULT 9
US-09-784-674-471/c
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Best Local Similarity
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                                                                                                                                                 GENERAL
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FILING DATE: 10-FEB-1998
ATTORNEY/AGENT INFORMATION:
NAME: Choi, Wendy A.
REGISTRATION NUMBER: 36,697
REFERENCE/DOCKET NUMBER: 10971
probe sequences NUMBER OF SEQUENCES: 1165
                                 APPLICANT: Shannon, Karen W.
Wolber, Paul K.
Delenstarr, Glenda C.
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                         ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                             19 ACTGCATTTACCA 7
                                                                                                                                                 INFORMATION:
                                                                                                                                                                                                                                                                                              2 ACTGCACTTNCCA 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 650-852-8063
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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Company M/S 20BO
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                                                                          Webb, Peter G
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Webb, Peter G.
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                                                                                                                                                                                                                                                                                                                                                                 Score 10.4;
Pred. No. 1
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RESULT 10
US-09-784-674-472/c
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INFORMATION FOR SEQ ID NO: 471:
SEQUENCE CHARACTERISTICS:
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COUNTRY: USA
ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HYPOTHETICAL: NO ANTI-SENSE: NO
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ADDRESSEE: Records Manager, Legal Department, Hewlett-Packard Company M/S 20BO.
                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 1165
                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Methods for evaluating oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2 ACTGCACTTNCCA 14
                                                                                                                       STATE: CA
                                                                                                                                              CITY: Palo Alto
                                                                                                                                                            ADDRESSEE: Records Manager, Legal Department, Hewlett-Packard
Company M/S 20BO
STREET: 3000 Hanover Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 09/021,701 FILING DATE: 10-FEB-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                             Application US/09784674 o. US20030054346A1
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                                                                                                                                                                                                                                                                                                      Delenstarr, Glenda C. Webb, Peter G. Kincaid, Robert H.
                                                                                                                                                                                                                                                                                                                                                                   Shannon, Karen W. Wolber, Paul K.
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84.6%;
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TELECOMMUNICATION INFORMATION:

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US-09-784-674-473/c
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: cDNA
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 10971464-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-236-2386
                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/784,674 FILING DATE: 15-Feb-2001 CLASSIFICATION: NO. US20030054346A1 available PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/021,701
FILING DATE: 10-FEB-1998
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                                                             APPLICATION NUMBER: 09/021,701
FILING DATE: 10-FEB-1998
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kincaid, Robert H.
TITLE OF INVENTION: Methods for evaluating oligonucleotide
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Company M/S 20B0
STREET: 3000 Hanover Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Choi, Wendy A. REGISTRATION NUMBER: 36,697
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/09/784,674 FILING DATE: 15-Feb-2001
                     NAME: Choi, Wendy A. REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                  COUNTRY: UZIP: 94304
                                                                                                                                                                                                                                                                                                                                                                      CITY: Palo Alto
STATE: CA
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REFERENCE/DOCKET NUMBER: 10971464-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/09784674
o. US20030054346A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Shannon, Karen
Wolber, Paul K.
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Webb, Peter G.
                                                                                                                                                                                                                                                                                                                                                     USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 10.4; DB 9
Pred. No. 1e+04;
                       36,697
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Legal Department, Hewlett-Packard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 9; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2
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US-09-784-674-474/c
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Best Local Similarity
Matches 11; Conser
                                                                                                                                TELEPHONE: 650-236-2386
TELEFAX: 650-852-8063
INFORMATION FOR SEQ ID NO: 474:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/784,674
                                                                                                                                                                                                                                              APPLICATION NUMBER: 09/021,701
FILING DATE: 10-FEB-1998
ATTORNEY/AGENT INFORMATION:
NAME: Choi, Wendy A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
TOPOLOGY: linear MOLECULE TYPE: cDNA HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
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                                                                                                                SEQUENCE CHARACTERISTICS
                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kincaid, Robert H. TITLE OF INVENTION: Methods for evaluating oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Shannon, Karen W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: cDNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16 ACTGCATTTACCA 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 ACTGCACTTNCCA 14
                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 36,
                                                                                                                                                                                                                                                                                                                                           FILING DATE: 15-Feb-2001 CLASSIFICATION: No. US20030054346A1 available
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Records Manager, Company M/S 20BO
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 650-852-8063
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 3000 Hanover Street
                                                                                          LENGTH: 20 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE:
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Webb, Peter G.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      USA
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                                                                                                                                                                                                                             36,697
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ANTI-SENSE: NO

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US-09-784-674-475/c
US-09-784-674-475/c
Sequence 475, Application US/09784674
Publication No. US20030054346A1
GENERAL INFORMATION:
                                                   В
                                                                                                                                                                                             SEQUENCE DESCRIPTION: SEQ ID NO: 475: US-09-784-674-475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; SEQUENCE DESCRIPTION: SEQ ID NO: US-09-784-674-474
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                                                                                                                    Query Match
Best Local S
Matches 11
                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 09/021,701
FILING DATE: 10-FEB-1998
ATTORNEY/AGENT INFORMATION:
NAME: Choi, Wendy A.
RECISTRATION NUMBER: 36,697
REFERENCE/DOCKET NUMBER: 1097146.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-236-2386
TELEFAX: 650-852-8063.
INFORMATION FOR SEQ ID NO: 475:
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Best Local Similarity
                                                                                                                                                                                                                                HYPOTHETICAL: NO ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                               TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE:.Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                   14
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                                                                                   2 ACTGCACTTNCCA 14
                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                 LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/784,674 FILING DATE: 15-Feb-2001 CLASSIFICATION: NO. US20030054346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Records Manager,
Company M/S 20B0
STREET: 3000 Hanover Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: CA
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Wolber, Paul
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Webb, Peter G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA
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                                                                                                                                     74.3%;
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                                                                                                                      0,
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                                                                                                                                   Score 10.4;
Pred. No. 1
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                                                                                                                    Mismatches
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                                                                                                                                     4; DB 9;
le+04;
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                                                                                                                                                      Length 20;
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                                                                                                                                     RESULT 15
US-09-887-145-27
US-09-887-145-27
; Sequence 27, Application US/09887145
; Publication No. US20030082139A1
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US-09-784-674-476/c
                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 11; Conserv
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                                                                                                                                   GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 36,697
REFERENCE/DOCKET NUMBER: 10:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-236-236
TELEFAX: 650-852-8063
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/784,674
FILING DATE: 15-Feb-2001
CLASSIFICATION: NO. US20030054346A1 available
PRIOR APPLICATION DATA:
       CORRESPONDENCE ADDRESS:
ADDRESSEE: David Prashker,
STREET: P.O. Box, 5387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                APPLICANT: Kim, Seur TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS
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                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE DESCRIPTION: SEQ ID NO: 476:
                                                                                                                                                                                                                                                                                                                                                                                                                                            ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            HYPOTHETICAL: NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Methods for evaluating oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                               13 ACTGCATTTACCA 1
                                                                                                                                                                                                                                                                                                 2 ACTGCACTTNCCA 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 09/021,701 FILING DATE: 10-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn.Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: CA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 3000 Hanover Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Records Manager, Legal Department, Hewlett-Packard Company M/S 20BO .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Choi, Wendy A.
                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shannon, Karen W. Wolber, Paul K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kincaid,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Delenstarr, Glenda C
Webb, Peter G.
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                                                                                                                  Seung U
                                                                                                                                                                                                                                                                                                                                                     74.3%;
                                                                             cell and continuous
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                                                                                                  Immortalized human microglia
                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                   Score 10.4; DB Pred. No. 1e+04;
                                                                                                                                                                                                                                                                                                                                      Mismatches
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9; 2

Length 20; Indels

0,

Gaps

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CITY: Magnolia
STATE: Massachusetts
COUNTRY: USA
2 IP: 01930
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.40 Mb storage
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.40 Mb storage
COMPUTER SEAL PC
COMPUTER: Dell PC
COMPUTER: DESCRIPTION LATA
AFDILATION UNBER: US/09/887,145
FILING DATE: 22-JUN-201
CLASSIFICATION UNBER: US/09/887,145
FILING DATE: 22-JUN-201
CLASSIFICATION UNBER: UBC-002
TELECOMMUTICATION UNBER: UBC-002
TELECOMMUTICATION UNBER: UBC-002
TELECOMMUTICATION UNFORMATION:
TELEPHONE: (978) 525-3794
INFORMATION TELEPHONE: (978) 525-3794
INFORMATION: SEQUENCE GLARACTERISTICS:
EQUENCE GLARACTERISTICS:
TYPE: INCLEACE acid
STRANDEDNESS: single
SEQUENCE DESCRIPTION: SEQ ID NO: 27:
US-09-887-145-27

OUS-09-887-145-27

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Title:
Perfect score:
Sequence:
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                         Database :
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Gapop 10.0 , Gapext 1.0
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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FEATURES	COMMENT	JOURNAL	TITLE	REFERENCE AUTHORS	ORGANISM	VERSION KEYWORDS	LOCUS DEFINITION ACCESSION
Institute of Medical Science, University of Tokyo 4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan Email: Synzukieims.u-tokyo.ac.jp Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano ,S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997). Location/Qualifiers	Contact: Yutaka Suzuki	mapping of mRNA start sites EMBO Rep. 2 (5), 388-393 (2001) 21272072	, H., Ota, T., Isogai, T., Tanaka, T., Morishita, S., Okubo, K., Sakaki, Y., Nakamura, Y., Suyama, A. and Sugano, S. Diverse transcriptional initiation revealed by fine, large-scale	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  1 (bases 1 to 50) Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J., Sese,J., Hata	Homo sapiens	AU105465.1 GI:13554986	AU105465 50 bp mRNA linear EST 30-AUG-2001 AU105465 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone HRC09846, mRNA sequence.  AU105465

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RESULT 3
AZ936711
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African clawed frog.
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Contact: Tadasu Shin-i
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                                                                                                                                                                                                                                                                                                                                                                                               Email: tshini@genes.nig.ac.jp
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Center For Genetic Resource Information
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kitayama,A.,
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Amphibia; Batrachia; Anura; Me
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                                                                                                                                                                                                 /note="Vector: pBSRN3; Site_1: NotI; Site_2: EcoRI; cDNAs were oligo-dT primed and directionally cloned. Staging according to Nieuwkoop and Faber. Library is subtracted and was constructed by N. Garrett and A.M. Zorn, (Wellcome/CRC Institute). "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /db_xref="taxon:9606"
/clone="HRC09846"
./clone="HRC09846"
./clone="bif="Sugano Homo sapiens cDNA library"
./clone="bifferential display comparison of untreated note="bifferential display comparison of untreated dimethylfumarate treated U937 cells"
a 12 c 14 g 12 t
                                                                                                                                                                                                                                                                                                                                          /db_xref="taxon:8355"
/clone="XL084o15"
                                                                                                                                                                                                                                                                                    /dev_stage="stage 25"
                                                                                                                                                                                                                                                                                                   /tissue_type="whole embryo"
                                                                                                                                                                                                                                                                                                                 Library
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Mesobatrachia; Pipoidea
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6.9e+04;
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           GSS
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Tel: 801 585 5606

Fax: 801 585 7177

Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.9

Plate: 0193 row: I column: 03

Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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University of Utah
Rm. 308, Biomedical Polymers Research
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                          was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWHD42 (gi14732114|gb)AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (female) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M0193103"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (http://www.jax.org/resources/documents/dnares/).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone_lib="Mouse 10kb plasmid UUGC2M library"
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91.7%;
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Pred. No.
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Mismatches 1;
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RESULT 5
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Contact: Nori Satoh
Department of Zoology
                                                                            Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genet
                                     Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0013 row: P column: 2:
                                                                                                                                                                   Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                         Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Welss,R.

Mouse whole genome scaffolding with paired end reads from 10kb
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1 (bases 1 to 44)
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Seq primer: CACACAGGAAACAGCTATGACC Class: plasmid ends
                                                                                                                                                                                                                            plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 22)
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AZ309394.1 GI:10350335
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clone UUGC1M0013P22 R, DNA sequence.
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/clone_lib="Nori Satch unpublished cDNA library,
/clone_lib="Nori Satch unpublished cDNA library,
/tissue_type="whole animal"
/dev_stage="larva"
/note="Vector: pBluescript SK"
9 c 11 g 15 t
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/db_xref="taxon:7719"
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91.7%;
                                                                                                                                                Polymers Research Bldg.,
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olumn: 22
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RESULT 6
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                                                                                                                                                                 University of Utah
Rm. 308, Biomedical Polymers Research
                                                                                                                                                                                                                                                                                                                                                                  1 (bases 1 to 29)
Dunn,D., Aoyagi,A., Barber,M.,
Islam,H., Longacre,S., Mahmoud
                  Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0060 row: I column: 03
Seg primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                             ,M., Rose,M., Rose,R.,
and Wright,D.,Weiss,R.
Mouse whole genome scal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AZ801886 29 bp DNA linear GSS 16 2M0060103R Mouse 10kb plasmid UUGClM library Mus musculus clone UUGC2M0060103 R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Location/Qualifiers
Class: plasmid ends
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gil4732114)gblAF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E coll XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              was blunt end-repaired with T4 DNA polymerase and T4 polynucleotides kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     was hydrodynamically sheared by repeated passage through a 0.005\ \mathrm{inch} orifice at constant velocity. The sheared DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0013P22"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (http://www.jax.org/resources/documents/dnares/). The DNA
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84.6%;
                                                                                                                                                                                                                                                                                                           scaffolding with paired end reads from 10kb
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VERSION
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                                                                 University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Res
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Bukaryota; Metazoa; Chordata; Sciurognathi; Muridae; Murinae; Mus
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 41)
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   Class: plasmid ends
                                     Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                            Tel: 801 585 5606 Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AZ663066
AZ663066.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AZ663066 41 bp DNA linear 100542D06R Mouse 10kb plasmid UUGC1M library Musclone UUGC1M0542D06 R, DNA sequence.
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWDA2 (gil4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and
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/note="Vector: PWD42nv; Purified genomic DNA from M.
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/clone="UUGC2M0060I03"
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84.6%;
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Pred. No. 1.3e+05;
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                                                                                                                                                                                                                                                                                                                                    Contact: Yutaka Suzuki
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                                                                                                                                                                                                                    Email:
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/organism="Homo sapiens"
/db_xref="taxon:9606"
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1 (bases 1 to 50)
Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J., Sese,J., Ha,H., Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo,K., Sakaki,Y., Nakamura,Y., Suyama,A. and Sugano,S.
Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
EMBO Rep. 2 (5), 388-393 (2001)
                           Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano,S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
                                                                                                                                                                Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AU102646 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone CAS05488, mRNA sequence.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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    Location/Qualifiers
                                                                                                                                    ysuzuki@ims.u-tokyo.ac.jp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."
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Location/Qualifiers
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/strain="C57BL/6J"
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/note="Vector: PWD42nv; Purified genomic DNA from M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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AZ470763.1 GI:10628888
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34 bp DNA linear GSS 04-OCT-200
1M0285F10F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0285F10 F, DNA sequence.
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: ddunn@genetics.utah.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and Wright,D.,Weiss,R.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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/clone_lib="Sugano Homo sapiens cDNA library"
/note="Differential display comparison of untreated
dimethylfumarate treated U937 cells"
a 7 c 16 g 8 t
                adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gil4732114|gb)AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated
                                                                                                             was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                      /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                               (http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                                                                                                                                                                        Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                   /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                          /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                          /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
                                                                                                                                                                                                                                                                                                                     /sex="Male"
                                                                                                                                                                                                                                                                                                                                                       /clone="UUGC1M0285F10"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   74.3%;
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0; Mismatches
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  insert adaptors and
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TITLE
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Submitted (17-JUN-2002) Weisshaar B., Max-Planck-Institut fuer zuechtungsforschung, Carl-von-Linne-Weg 10, Koeln, 50829, Germany This sequence is recovered from the left border of the T-DNA, It indicates an insertion close to or within gene At2948150. The sequences are generated at the MPI for Plant Breeding Research in the context of the GABI-Kat project. GABI-Kat is part of the German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       genomic survey sequence
AL772108
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Arabidopsis thaliana T-DNA flanking
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         availability can be found at:
http://www.mpiz-koeln.mpg.de/GABI-Kat/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Strizhov, N., Li, Y., Rosso, M., Viehoever, P., Dekker, K., Saedler, H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Spermatophyta; Magnollophyta; eudicotyledons; core et
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                                                                                                                                                                                                                  /Clone="GK-186B11-014622"
/clone_lib="Arabidopsis thaliana T-DNA insertion lines"
/clone_lib="Arabidopsis thaliana T-DNA insertion lines"
/clone_"PCR was performed on DNA from Arabidopsis thaliana
plants (T1) which were transformed with the T-DNA from
vector pAC161. The lines contain one or more T-DNA
insertions. The DNA fragment(s) resulting from the PCR
were directly sequenced to determine the genomic sequence
                                                                                                                                      similarity
                                                                                                                                                                                      flanking the insertion.
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                                                                                      for submission. T-DNA
                                                                                                                                 he insertion. Sequences displaying significant to the A. thalaina nuclear genome sequence wea
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Pred. No. 2.2e+05;
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71.48;

Score 10;

DΒ 17;

Length 46;

Indels

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Gaps

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Pipoidea;

Euteleostomi;

Pipidae;

Shin-i, T. and Kohara

linear

ear EST a library

06-DEC-2001 Xenopus

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BASE COUNT
ORIGIN
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clone UUGCIM0269K14 F, DNA sequence.
AZ462142
AZ462142.1 GI:10620267
GSS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Resonance Rm. 308, Biomedical Polymers Rm. 308, Biomedical P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         High quality sequence stop: 47.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0269 row: K column: 14
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mouse whole genome scaffolding with paired end
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 (bases 1 to 47)
Dunn, D., Aoyagi, A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus
Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mammalia; Eutheria;
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801 585 7177
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                                                                                                                                                                                     (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored work of the control work of the control vector DNA.
                                                                                                a
                                                                                        adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."

1 c 25 g 9 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone="UUGC1M0269K14"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          'lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    90.9%;
                   71.48;
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Rodentia;
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                   Score
                   10;
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             Length 47;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           from the Jackson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         source
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C20876
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           Okubo, K.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BJ034300 NIBB Mochii normalized Xenopus neurula laevis cDNA clone XL028b09 5', mRNA sequence.
                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                   Homo sapiens
                                                                                                                                                                                                C20876.1
                                                                                                                                                                                                                                                                                                         C20876
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Kitayama, A., Terasaka, C., Mochii, M., Ueno, N.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Tadasu Shin-i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Expressed genes in X. laevis embryo Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Xenopus laevis
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                                                                                                                                               numan.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Amphibia; Batrachia; Anura; Mesobatrachia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    African clawed frog.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             tshini@genes.nig.ac.jp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         according to Nieuwkoop and Faber. Library is subtracted and was constructed by N. Garrett and A.M. Zorn, (Wellcome/CRC Institute). "" 2 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /tissue_type="whole embryo"
/dev_stage="stage 15"
/note="Vector: pBSRN3; Site_1: NotI; Site_2: EcoRI; cDNAs
were oligo-dT primed and directionally cloned. Staging
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone="XL028b09"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /db_xref="taxon:8355"
                                      1 to 49)
                                                                                                                                                                                                GI:1621986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone_lib="NIBB Mochii normalized
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ∕organism="Xenopus laevis"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Metazoa; Chordata; Craniata; Vertebrata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GI:17391841
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                                                                                                                                                                                                                                                                             Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Pred. No. tive 0; Mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                          26
                                                                                                                                                                                                                                                                             adult
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 10; DB 13;
Pred. No. 2.4e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                          (K.Okubo)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                                                                                                                                                                                                                                                      mRNA
                                                                                                                                                                                                                                                                          Homo sapiens cDNA
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Xenopus neurula

Length 48;

0;

Gaps

0;

linear

EST 23-OCT-1996

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JOURNAL COMMENT
                                                                                                                                                                                                                                                                                                                    REFERENCE
AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                               ACCESSION
VERSION
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AZ481615
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ORIGIN
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Best Local (
                                                                                                                                                                                                      JOURNAL
                                                                                                                                                                                                                                          TITLE
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                                                                                                                                            University of Utah Guiversity of Utah
                                                                                                                                                                                                                                                                                                            AZ481615 25 bp DNA linear GSS V4-ULI-ZVV
1M0306A03F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0306A03 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: kousaku@imcb.osaka-u.ac.jp
Human Gene Signature, 3'-directed cDNA sequence. We are not
submitting the same cDNA sequence redundantly to DDBJ since 1993.
For the abundance information of clones with this sequence in this
library and as well as in other 3'-directed libraries, see '
http://www.imcb.osaka-u.ac.jp/bodymap'. The sequences of the clones
represented by this GS sequences is also found there.
Location/Qualifiers
                                                                                                                                                                                                                                                          Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   9
                   Plate: 0306
                                                                       Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                              GSS
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Seq primer: CGTTGTAAAACGACGGCCAGT
                                Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
                                                                                                                                                                                                                                         Mouse whole genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                 AZ481615.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Osaka University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Okubo, K
                                                                                                                                                                                                                         plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                            house mouse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Institute for Molecular and Cellular Biol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                 308,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                               USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="Organ: blood; Vector: 1-gt-11; Site_1: Eco-RI; Monocytes were prepared from blood by ficoll-hypaque, percoll and T cell rosetting purification steps (purity: 96 %). mRNA was prepared from activated monocytes from a patient with rheumatoid arthritis. mRNA was reverse transcribed with MuLV. Using Eco-RI linkers cDNA was cloned into 1-gt-11 vector arms. The cDNA library was screened by differential hybidization using radioactivel, marked ss-cDNA from activated and non-activated
                                                                                                                             Biomedical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 monocytes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone_lib="Human adult (K.Okubo)"
/dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                               GI:10642680
                   row: A column:
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                                                                                                                                                                 Genome
                                                                                                                                                                                     Weiss
                                                                                                                                                                                                                                     scaffolding with paired
                                                                                                                           Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 10; DB 14;
Pred. No. 2.4e+05;
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                                  Std Error: 0.00
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RESULT 15
HSTEA0309
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ORIGIN
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                                                                                                                                                                                                                                                         MEDLINE
PUBMED
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Best Local :
                                                                                                                                                                  JOURNAL
                                                                                                                                                                                                             AUTHORS
                                                                                                                                                                                                                                                                                                       JOURNAL
                                                                                                                                                                                   2 (bases 1 to 26)
Antonarakis, S.E.
Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26 bp H.sapiens trapped exon SNSA0309, x85346
                                                                                                                                                             Submitted (14-MAR-1995) S.E.
                                                                                                                                                                                                                                                                                                         Hum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GSS; chromosome 21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 CACTGCACTINCCA 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Class: plasmid ends
                                                                                                                                         Univ.
                                                                                                                                                                                                                                                                                                                               syndrome
                                                                                                                                                                                                                                                                                                                                                                        Lucente,D., Chen,H.M., Shea,D., Samec,S.N., Rutter,M., Chrast,R.,
ROSsier,C., Buckler,A., Antonarakis,S.E. and McCormick,M.K.
                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostómi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 26)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   High quality sequence stop: 25
                                                                                                                                                                                                                                                                                   96090248
                                                                                                                                                                                                                                                                                                                                                Localization of 102 exons to a 2.5 Mb region involved
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                                                                                                                                                                                                                                                                                                    Mol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                 and Cantonal Hospital of
211 Geneva, SWITZERLAND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gil473214) pblnF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and
/db_xref="taxon:9606"
/chromosome="21"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA
                                             /organism="Homo sapiens"
                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                       Genet. 4 (8), 1305-1311 (1995)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="Vector:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /db_xref="taxon:10090"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone="UUGC1M0306A03"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     70.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 9.8; DB 17;
Pred. No. 2.7e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                             Antonarakis, Div.
                                                                                                                                         Geneva, CMU,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         genomic survey sequence
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                                                                                                                                       1 Rue Michel-Servet,
                                                                                                                                                               of Medical Genetics,
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Anse count 2 a 11 c 4 g 9 t
ORIGIN

Query Match
Best Local Similarity 78.6%; Score 9.8; DB 17; Length 26;
Best Local Similarity 78.6%; Pred. No. 2.7e+05;
Matches 11; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

Qy 1 CACTGCACTTNCCA 14
Db 12 CCCTGCACTGTCCA 25
Search completed: June 21, 2003, 23:59:01
Job time : 610.901 secs
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Post-processing; Minimum Match
Maximum Match
Listing first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Database :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Minimum DB seq length: 0 Maximum DB seq length: 50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Scoring table:
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Run on:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OM nucleic -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nucleic search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        June 21, 2003, 21:23:11; Search time 281.094 Seconds (without alignments) 1449.478 Million cell updates/sec
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14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       '2054640 seqs, 14551402878 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IDENTITY_NUC
Gapop 10.0 , Gapext 1.0
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gb_ph:*
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em_pat:*
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REFERENCE AUTHORS TITLE	KEYWORDS SOURCE ORGANISM	RESULT 1 AX190063 LOCUS DEFINITION ACCESSION VERSION		C 44 C 45	C 42	C 40			C 35	•	C 32		C 28	26 27	25 \$	23	21 22 :	20	18 19	16 17	c 15		12	11			7 O			2 1-	Result	
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1 (bases 1 to 15)
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Copelstein, B., Kinzler, K.W., Zhang, L. and Zhou, W.
Gene expression profiles in normal and cancer cells
Patent: US 6333152-A 48 25-DEC-2001;
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Nucleic acids containing single nucleotide
methods of use thereof
Patent: WO 0147942-A 244 05-JUL-2001;
Curagen Corporation (US)
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/db_xref="taxon:9606"
/note="2 of 2 allelic variants
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Cargill,M., Ireland,J.S. and Lander,E.S.
Human single nucleotide polymorphisms
Patent: WO 0166800-A 1670 13-SEP-2001;
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH
1 CCAGGTGTAGNCC 13
                                                                                                                               l (bases 1 to 15)
Stinchcomb, D.T., Jarvis, T. and McSwiggen, J.
Nucleic acid based inhibition of CD40
Patent: US 6194150-A 820 27-FEB-2001;
                                                                                                                                                                                      Unknown
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Sequence
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                                                                                                                                                                                Submitted (04-AUG-2001) Akira Horii, Tohoku University School of Medicine, Molecular Pathology; 2-1 Seiryomachi, Aoba-ku, Sendai, Miyagi 980-8575, Japan (E-mail:horii@mail.cc.tohoku.ac.jp, Tel:81-22-717-8042, Fax:81-22-717-8047)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                     and Soeda,E.

A BAC-based STS-content map spanning a 35-Mb region of chromosome 1p35-p36
Genomics 74 (1), 55-70 (2001)
                                                                                                                                                                                                                                                                                                                                                                    Chen,Y.Z., Hayashi,Y., Wu,J.G., Takaoka,E., Maekawa,K., Watanabe,N., Inazawa,J., Hosoda,F., Arai,Y., Mizushima,H. Morohashi,A., Ohira,M., Nakagawara,A., Liu,S., Hoshi,M.,
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                                                                                                                                                Hirano,T., Nakane,S., Mizoshita,K., Yamakuchi,H.,
Inoue-Murayama,M., Watanabe,T., Barendse,W. and Sugimoto,Y.
Characterization of 42 highly polymorphic bovine microsatellite,
                                                                                                                                                                                                                       Inoue, M., Hirano, T., Nakane, S., Watanabe, T., Takeda, H., Yamakuchi, H., Morita, M., Barendse, W. and Sugimoto, Y. Five bovine polymorphic dinucleotide microsatellite loci DIK010, DIK015, DIK016 and DIK020)
Anim. Genet. 26 (6), 447-448 (1995)
           Unpublished
                                                   Inoue,M., Watanabe,T., Hirano,T., Yamakuchi,H., Tsukazawa,H., Watanabe,E., Morita,M. and Sugimoto,Y.
Isolation of microsatellites from Japanese black cattle (Wagyu)
                                                                                                            Anim. Genet.
97083737
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Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pec
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Bonthron, D. and Markham, A.F.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae
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                                       their application
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/ob_xref="taxon:9606"
/note="Oligonucleotide Primer"
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DNA PCR a

antisense primer

Pecora;

Bovoidea;

(DIK008,

Euteleostomi;

linear

MAM 09-FEB-1999

(GB)

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MARKHAM ALEXANDER FRED

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Hominidae;

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PAT 16-SEP-2000

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JOURNAL
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AR099541
AR099541.1
                                          Sharpe,A.H., Borriello,F., Freeman,G.J. B7-1 and B7-2 polypeptides Patent: US 6218510-A 61 17-APR-2001;
                                                                                                                                AR146442
Sequence 61 from patent
AR146442
AR146442.1 GI:15109631
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Bennett, C. Frank. and Vickers, T.A.
Oligonucleotide compositions and methods
expression of B7 protein
Patent: US 6077833-A 68 20-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted (21-DEC-1994) Yoshikazu Sugimoto, Japan Live Technology Association, Shirakawa Institute of Animal G Nishigo Odakura, Nishishirakawa, Fukushima 961, Japan (E-mail:LDI03222@niftyserve.or.jp, Tel:0248-25-5641,
                                                                                                  Unknown
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/db_xref="taxon:9913"
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 organism="unknown" 6 c 5 g
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Query Match
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                                                                                                                                                             Li,X., Ohmen,J.D., Reed,D.R., Ross,D. & Gene and sequence variation associated compounds and other sweeteners Patent: WO 0183749-A 238 08-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 20)
Bennett, C. Frank. and Vickers, T.A.
Oligonucleotide compositions and methods
expression of B7 protein
Patent: US 6319906-A 68 20-NOV-2001;
                                                                                                                                                                                                    Bachmanov,A.A., Beauchamp,G.K., Chatterjee,A., de Jong,P.J., Li,S., Li,X., Ohmen,J.D., Reed,D.R., Ross,D. and Tordoff,M.G.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
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1 (bases 1 to 22)
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                                                                               Patent: WO 0146391-A 209 28-JUN-2001,
Plant Bioscience Limited (GB)
Location/Qualifiers
1, 22
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Sequence 209 from Patent W00146391.
AX180631 GI:15132517
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Submitted (04-AUG-2001) Akira Horii, Tohoku University School of Medicine, Molecular Pathology; 2-1 Seiryomachi, Aoba-ku, Sendai, Miyagi 980-8575, Japan (E-mail:horii@mail.cc.tohoku.ac.jp, Tel:81-22-717-8042, Fax:81-22-717-8047)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A BAC-based STS-content map spanning a 35-Mb region of human chromosome 1p35-p36 Genomics 74 (1), 55-70 (2001) 21269192
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    synthetic construct DNA synthetic construct artificial sequences.
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sts-R272C15F obtained from clones B359F10, B272C15,
B91D18, Human BAC library RPCI-11"
a 4 c 6 g 6 t
             /organism="synthetic construct"
/db_xref="taxon:32630"
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AAAIJOOJ	AAX15861	AAA53225	AAX98230	AAQ94196	AAQ94195	AAV17107	AAX89404		AAT33447 .	AAI30183 ·	AAT97150	ABK87566	AAV55428	AAT62986	AAT65516	AAX34310	ABK66989	AAT34014	AAS09815 .	AAA59964	AAV21625	AAT11220	ABL45384	ABL44107	AAS97628	AAS96640	AAD34980	AAF32871	314	AAA91878	AAV48029	AAT01071	S	60	AAA52824
PCK PIIMEI USEG CO	R primer used t	Human G-protein co	Nucleotide sequenc	Transmembrane olig	smembrane oli	erine/threor	PCR pr	Human pax protein	MAb chMint5 light	Human single nucle	Primer S1 for ob p	Matrix metalloprot	Primer 35-5' for c	c-mpl receptor ago	Oligonucleotidé 35	Human oestrogen re	Human gene speçifi	obacteria 168	Oat Beta-amyrin sy	imer for th	þţ	Internal control p	chromosome	hromosome	gene-	revers	ATHER gene		lec	se primer	7 - 1	Mouse B7-1 (altern	nucleotide u	chrom	Human CD44 antisen

## ALIGNMENTS

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RESULT 1
AAH90362
Key
variation
       Shimkets RA,
                                         27-DEC-1999;
                                                         27-DEC-2000; 2000WO-US35387
                                                                           05-JUL-2001.
                                                                                             WO200147942-A2
                                                                                                                                                       Homo sapiens
                                                                                                                                                                       Human; single nucleotide polymorphism; SNP; chromosome 19;
detection; identification; gene therapy; genetic disorder;
                                                                                                                                                                                                  Human clone cg43982025 SNP site, SEQ ID NO:242
                                                                                                                                                                                                                    08-OCT-2001
                                                                                                                                                                                                                                     AAH90362;
                                                                                                                                                                                                                                                      AAH90362 standard; cDNA; 50
                       (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                  (first entry)
      Leach M;
                                         99US-0472865
                                                                                                                             replace(25..26,GCC)
                                                                                                            /standard_name= "single nucleotide polymorphism"
                                                                                                                                                                       genetic disorder;
                                                                                                                                                                         SS.
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RESULT 2
AAH90364
XXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             contain silent SNPS. Sequences 569 to 580 (AAH90689-AAH90700) are consecutive pairs of nucleotides containing SNPs which result in changes in the corresponding amino acid sequences (AAG64751-AAG64762). The SNPS in sequences 569 to 574 (AAH90689-AAH90694) lead to conservative amino acid changes, while those in sequences 575 to 578 (AAH90698) result in non-conservative changes. The SNP in sequences 579 and 580 (AAH90699-AAH90700) generates a frameshift mutation. The invention also relates to a method of detecting a polymorphic site in a nucleic acid and a method of determining the relatedness of two nucleic acids. It also encompasses peptides containing polymorphic sites, antibodies ralsed against such peptides, and a method of detecting polymorphic proteins/ capations to a individual having, suspected of having, or at risk of developing a pathological condition due to the presence of a sequence polymorphism. Such treatment would comprise administration of the wild-type nucleic acid sequence. Antibodies ralsed against polymorphic proteins/ peptides can also be used in the treatment of such individuals.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequences AAH90121-AAH90700 represent 580 human cDNA sequences which contain single nucleotide polymorphisms (SNPs). Sequences 1 to 568 (AAH90121-AAH90688) are consecutive pairs of nucleotides which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New polynucleotides containing single nucleotide polymorphisms, for detecting the presence of polymorphism, detecting a polymorphic site, and treating a patient suffering from a pathology ascribed to the
                 Shimkets RA,
                                               (CURA-) CURAGEN CORP.
                                                                                 27-DEC-1999;
                                                                                                                  27-DEC-2000; 2000WO-US35387
                                                                                                                                                                                       W0200147942-A2
                                                                                                                                                                                                                                                         variation
                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                          detection; identification;
                                                                                                                                                                                                                                                                                                                                                        Human; single nucleotide polymorphism; SNP; chromosome 19;
                                                                                                                                                                                                                                                                                                                                                                                                                           08-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAH90364;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAH90364 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 50 BP; 9 A; 13 C; 21 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Page 87; 295pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-425617/45
                                                                                                                                                                                                                                                                                                                                                                                          clone cg43982025 SNP site, SEQ ID NO:244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 CCAGGTGTAGNCCA 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CCAGGTGCAGGCCA 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
               Leach M;
                                                                                 99US-0472865
                                                                                                                                                                                                                                     Location/Qualifiers replace(25..26,CCA)
/*tag= a
                                                                                                                                                                                                                    /standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 81.4%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                50
                                                                                                                                                                                                                                                                                                                                        gene therapy; genetic disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 11.4; DB 22; Pred. No. 3.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
                                                                                                                                                                                                                                                                                                                                          SS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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RESULT 3
AAX30994/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CC contain silent SNPs. Sequences 569 to 580 (AAH90689-AAH90700) are CC consecutive pairs of nucleotides containing SNPs which result in changes in the corresponding amino acid sequences (AAG64751-AAG64762). The SNPs CC in sequences 569 to 574 (AAH90689-AAH90694) lead to conservative amino CC acid changes, while those in sequences 575 to 578 (AAH90698) CC result in non-conservative changes. The SNP in sequences 599 and 580 (CC (AAH90699-AAH90700) generates a frameshift mutation. The invention also CC relates to a method of detecting a polymorphic site in a nucleic acid and CC a method of determining the relatedness of two nucleic acids. It also CC encompasses peptides containing polymorphic sites, antibodies raised CC against such peptides, and a method of detecting polymorphic proteins/ CC peptides using the antibodies. The nucleic acids are useful for genee therapy of an individual having, suspected of having, or at risk of CC developing a pathological condition due to the presence of a sequence CC polymorphicm. Such treatment would comprise administration of the CC wild-type nucleic acid sequence. Antibodies raised against polymorphic proteins conditions are useful to sequence wild-type nucleic acid sequence. Antibodies raised against polymorphic conditions are useful to sequence wild-type nucleic acid sequence. Antibodies raised against polymorphic conditions are useful to sequence administration of the conditions are useful to sequence wild-type nucleic acid sequence. Antibodies raised against polymorphic conditions are useful to sequence administration of the conditions are useful to sequence and sequence and sequence and individuals.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequences AAH90121-AAH90700 represent 580 human cDNA sequences which contain single nucleotide polymorphisms (SNPs). Sequences I to 568 (AAH90121-AAH90688) are consecutive pairs of nucleotides which contain silent SNPs. Sequences 569 to 580 (AAH90689-AAH90700) are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New polynucleotides containing single nucleotide polymorphisms, for detecting the presence of polymorphism, detecting a polymorphic site, and treating a patient suffering from a pathology ascribed to the
                                                                                                                                                                                                                                                                                                                                                                                                  diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tag sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX30994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAX30994 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 87;
                                                                                                                                                                                                  21-MAY-1997;
                                                                                                                                                                                                                                        20-MAY-1998;
                                                                                                                                                                                                                                                                               26-NOV-1998.
                                                                                                                                                                                                                                                                                                                       W09853319-A2
                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                         Tag sequence; colorectal cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-425617/45
                                                                                                                                                        (UYJO ) UNIV JOHNS HOPKINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18 CCAGGTGCAGGCCA 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 CCAGGTGTAGNCCA 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                  prognosis; treatment;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                             of a transcript increased in colorectal cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                  97US-0047352
                                                                                                                                                                                                                                          98WO-US10277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             295pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         81.4%;
85:7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 11.4; DB 27
Pred. No. 3.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                     pancreatic cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                                                                                                                                                                                                         colon cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          50;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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0;

Use of isolated gene transcripts - useful for developing products for the diagnosis, prognosis and treatment of cancers, particularly colon and pancreatic cancer

WPI; 1999-070161/06

XW.

Vogelstein B;

Claim 2;

Page

24; 120pp;

English.

sequences of transcripts that

AAX30947-31815

Qy

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RESULT 4
ABK31947/c
ABK319
В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       differentially expressed in colorectal cancer, in pancreatic cancer, or in both. The tag sequences can be used to identify genes by matching the tag to a gen data base member, or by using the tag sequences as probes to isolate unidentified genes from cDNA libraries. The tag sequences can also be used in a method for diagnosing colon or pancreatic cancer in a sample suspected of being neeplastic. The method comprises comparing the level of at least one transcript in a first sample of a tissue to a second sample, where the first sample is a colonic tissue suspected of being neoplastic and the second sample is a normal human colonic tissue. The transcript is identified by a tag selected from AAX30947-31815. The methods of the invention can be used in the
                            that has the same sequence as a mRNA found in humans and is a SAGE (serial analysis of gene expression) tag comprising a single stranded probe containing at least loconsecutive nucleotides. SAGE tags, are diagnostic and prognostic markers of cancer, especially of the colon and pancreas. ABK31900-ABK32770 represent human colon and pancreatic cancer SAGE tags of the invention.
                                                                                                                                                           Disclosure; Column 15; 161pp;
                                                                                                                                                                                           New human nucleic acid diagnostic markers for
                                                                                                                                                                                                                                          WPI; 2002-153821/20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABK31947;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABK31947 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diagnosis, prognosis and treatment of cancer
                                                                                                                             The invention relates to an isolated, purified human nucleic acid
                                                                                                                                                                                                                                                                           Vogelstein B,
                                                                                                                                                                                                                                                                                                         (UYJO ) UNIV JOHNS HOPKINS
                                                                                                                                                                                                                                                                                                                                         20-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                       20-мау-1998;
                                                                                                                                                                                                                                                                                                                                                                                                        25-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                        US6333152-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cancer marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       serial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human colon cancer SAGE tag #48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-APR-20,02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                colon cancer; colorectal cancer; pancreatic cancer; SAGE tag; analysis of gene expression; diagnostic; prognostic; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ω
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ss.
                                                                                                                                                                                                                                                                           Kinzler
                                                                                                                                                                                                                                                                                                                                         98US-0081646
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
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                                                                                                                                                                                                                                                                           KW,
                                                                                                                                                                                         containing specific scancer, also derived
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15
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                                                                                                                                                                                                                                                                         Zhang
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Pred.
                                                                                                                                                           English
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                                                                                                                                                                                                                                                                           Zhou
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8e+03;
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                                                                                                                                                                                                                                                                         Σ
                                                                                                                                                                                                          SAGE
                                                                                                                                                                                           probes
                                                                                                                                                                                                        tags,
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                                                                                                                                                                                                          useful as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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Sequence 15

BP; 3

A; 5

C; 3

G; 4 T; 0 other

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14 CAGGTGTAGACC 3

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Qγ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best
                                                       Matches
                                                                               Best Local
                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                       The invention relates to the identification of nucleic acid molecules (AAII29513-AAI31314) from the human genome which include polymorphic sites which can predispose individuals to disease. Various genes from a number of individuals were resequenced and single nucleotide polymorphisms (SNPs) in these genes discovered. The method is useful for predicting the presence, absence or severity of a particular phenotype or disorder (e.g. diabetes) associated with a particular genotype. The nucleic acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleic acid molecules from the human genome which include polymorphic sites, useful in methods for predicting the presence, absence or severity of a particular phenotype or disorder (e.g. diabetes) associated with a particular genotype
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; resequence; genotype; disease
single nucleotide polymorphism; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-MAR-2000;
22-MAY-2000;
                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 132; 145pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    associated with a particular genotype
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-522952/57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cargill M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-MAR-2001; 2001WO-US07268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-SEP-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAI31182;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAI31182 standard; DNA; 31 BP
                                                                                                                                                                                                                                  containing the polymorphic sites may be useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (WHED)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           S
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                                                       11;
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                                                                             Similarity
  CAGGTGTAGNCC
                                                                                                                                                             3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ireland
                                                                                                                                                          BP;
                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000US-0187510
2000US-0206129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nucleotide polymorphism (SNP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                          8 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 replace(16,T)
/*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              genotype; disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                78.6%;
91.7%;
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                                                                                  91
                                                                                                          78.6%;
                                                                                                                                                          Ç;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BIOMEDICAL RES
                                                                               . 78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lander ES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English.
                                                                                                                                                       G; 6 T; 0
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Pred. No.
                                                                               Pred. No.
                                                                                                          Score 11;
                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SS
                                                                                                                                                          other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              forensic; paternity testing;
                                                                               6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5.8e+03;
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                                                                                                          DΒ
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                                                                                                                                                                                                                                     in
                                                                                                  Length 31,
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                                                                                                                                                                                                                                     forensics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                       Indels
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                                                                                                                                                                                                                                  paternity
                                                       Gaps
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밁 Q

15

CCAGGTGAAGTCC

Matches

11;

Conservative

0,

Mismatches

2;

Indels

0,

Gaps

0

1 CCAGGTGTAGNCC

13

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CC The present invention describes a novel enzymatic nucleic acid (ENA) CC having a hammerhead motif (HM) comprising: (i) at least 5 ribose CC residues; (ii) a 2'-C-allyl modification at position 4 of the ENA; (iii) CC at least ten 2'-O-methyl modifications; and (iv) a 3'-end modification. CC The ENA's can inhibit collagenase and stromelysin production in the CS synovial membrane of joints for the treatment or prevention of arthritis. CC particularly osteoarthritis or rheumatoid arthritis. The ENA's can also CC be used to treat antigen presenting cells of a donor to induce tolerance CC in a'recipient to an alloantigen of a donor. They can also be used for CC enhancing graft tolerance or for treating autoimmune disease, and for CC treating allergies and other inflammatory conditions. The ENA's can also CC be used in diagnosis. Ribozyme therapy impacts on the expression of CC stromelysin without introducing the non-specific effects upon gene CC expression which accompany treatment with retinoids and dexamethasone. CC The concentration of ribozyme required to affect a therapeutic treatment
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AAX65326/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-APR-1995;
02-MAY-1995;
04-MAY-1995;
07-JUL-1995;
07-JUL-1995;
07-AUG-1995;
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13-DEC-1994;
23-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-DEC-1994;
23-DEC-1994;
17-FEB-1995;
                                                                                                                                                                                                                                                                                                   Claim 10; Page 179;
                                                                                                                                                                                                                                                                                                                                treatment of auto-immune diseases
                                                                                                                                                                                                                                                                                                                                                       Enzymatic nucleic acid molecules having
                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1996-300653/30
                                                                                                                                                                                                                                                                                                                                                                                                                   Matulic-Adamic
                                                                                                                                                                                                                                                                                                                                                                                                                               Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-NOV-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-JUN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9618736-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       stromelysin; synovial rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX65326 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hammerhead ribozyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Arthritic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mouse
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (RIBO-)
                                                   lower than that required of
                                                                                                                                                                                                                                                                                                                                             the treatment of arthritis,
                                                                                                                                                                                                                                                                                                                                                                                                                                             ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  condition;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Gustofson J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hammerhead ribozyme target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                 Karpeisky A,
c J, Jarvis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            94US-0363253.
94US-0363254.
95US-0390850.
95US-0426124.
95US-0432874.
95US-0434509.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 95US-0000974
95US-0512861
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RNA;
                                                                                                                                                                                                                                                                                               307pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               graft tolerance;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hairpin ribozyme; human; rabbit; mouse; collagenase; membrane; joint; arthritis; osteoarthritis; autoimmune disease; allergy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15
                                                                                                                                                                                                                                                                                                                                                                                                                             McSwiggen J, Pavco
A, Modak A, Usman
                                                                                                                                                                                                                                                                                                                                                                                                                 Ή,
                                                                                                                                                                                                                                                                                                  English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВP
                                                                                                                                                                                                                                                                                                                                                                                                                Thompson
                                      antisense molecules, and is highly is used in the exemplification of
                                                                                                                                                                                                                                                                                                                                          s having a induction
                                                                                                                                                                                                                                                                                                                                                                                                        Pavco P, Stince.
Usman N, Burgin A;
"Fincott F;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID
                                                                                                                                                                                                                                                                                                                                          hammer-head motif - of graft tolerance of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             response;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NO:1958
                                                                                                                                                                                                                                                                                                                                                                                                                                          Stinchcomb DT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             target;
                                                                                                                                                                                                                                                                                                                                            or
                                                                                                                                                                                                                                                                                                                                                      used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cleavage;
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20 APR 1995; 02 MAY 1995; 04 MAY 1995; 07 JUL 1995; 07 JUL 1995;

95US-0000951 95US-0000974

95US-0434509

95US-0512861

95US-0426124 95US-0432874

95US-0390850 94US-0363253. 94US-0363254 94US-03

05-OCT-1995; 13-DEC-1994; 23-DEC-1994; 23-DEC-1994; 17-FEB-1995;

22-NOV-1995;

95WO-US15516

20-JUN-1996.

WO9618736-A2

SS

Arthritic condition; graft tolerance; immune response; target; cl hammerhead ribozyme; hairpin ribozyme; human; rabbit; mouse; coll stromelysin; synovial membrane; joint; arthritis; osteoarthritis; rheumatoid arthritis; autoimmune disease; allergy; inflammation;

collagenase; cleavage; Mouse B7-1 hammerhead ribozyme target

SEQ

IJ

NO:1959

AAX65327;

20-JUL-1999

(first entry)

AAX65327 standard; RNA; 15

ВP

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RESULT 7
ANA65327/c
ID ANA653
XX MOUSE
XX AITHRI
KW AITHRI
ANA9
PN 05-OCT
PR 05-OCT
PR 05-OCT
PR 13-DEC
PR 23-DEC
PR 13-DEC
PR 07-AUG
XX DIABLE
PT 10-ANAY
PR 07-AUG
XX Enzyma
PT for th
PT treatm
XX Enzyma
PT for th
PT treatm
XX Claim
XX The pr
CC Tabe pr
CC Tabe pr
CC Tabe or
CC Tabe use
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CC
                       The ENA's can inhibit collagenase and stromelysin production in the synovial membrane of joints for the treatment or prevention of arthritis, particularly osteoarthritis or heumatoid arthritis. The ENA's can also be used to treat antigen presenting cells of a donor to induce tolerance in a recipient to an alloantigen of a donor. They can also be used for enhancing graft tolerance or for treating autoimmune disease, and for treating allergies and other inflammatory conditions. The ENA's can also be used in diagnosis. Ribozyme therapy impacts on the expression of stromelysin without introducing the non-specific effects upon gene expression which accompany treatment with retinoids and dexamethasone. The concentration of ribozyme required to affect a therapeutic treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                    having a hammerhead motif (HM) comprising: (i) at least 5 ribose residues; (ii) a 2'-C-allyl modification at position 4 of the ENA; (iii at least ten 2'-O-methyl modifications; and (iv) a 3'-end modification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes a novel enzymatic nucleic
lower than that required of antisense
       molecules,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  acid
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Claim 10; Page

179; 307pp; English.

Enzymatic nucleic acid molecules having a hammer-head motif - used for the treatment of arthritis, induction of graft tolerance or treatment of auto-immune diseases

Beigelman L, Karpeisky A, N Matulic-Adamic J, Jarvis T,

McSwiggen J, Pavco A, Modak A, Usman s T, Thompson JD,

Wincott Ρ,

Burgin Stinchcomb

DΤ;

Pavco

~

Gustofson

1996-300653/30

(RIBO-) 07-AUG-1995;

RIBOZYME PHARM INC

Query Match Best Local Similarity

74.3%; 84.6%;

Score 10.4; Pred. No. 1.

DB 17;

Length 15

Sequence

15

ВP;

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4

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G

G;

3 U;

0

other;

present invention.

specific.

present

sequence

SSSS

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RESULT 8
ABL52004/c
ID ABL520
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                       The present invention describes an isolated polynucleotide (I) having a sequence (S1) comprising soluble carrier family 18 (vesicular monoamine), member 2 (SLC18A2) isogene selected from 49 isogenes with regions of a sequence (SS) of 40023 bp (see ABL51954), and defined by a corresponding set of polymorphisms whose locations and identities are given in the specification; or a sequence (S2) complementary to (S1). (I) has antioflammatory and neuroleptic activities, and can be used in gene therapy. Methods from the present invention can be used for haplotyping and genotyping the SLC18A2 gene in an individual SLC18A2 is also known and the motor of the present invention can be used in also known and senotyping the SLC18A2 gene in an individual SLC18A2 is also known and the motor of the present invention can be used in also known and the motor of the present invention can be used in also known and the motor of the present invention can be used in also known and the motor of the present invention can be used in the second of the present invention can be used in the second of the present invention can be used in the second of the present invention can be used in the second of the present invention can be used in the second of the present invention can be used in the second of the present invention can be used in the second of the present invention can be used in the second of the present invention can be used in the present invention can be used in the second of the present invention can be used in the present 
                         and genotyping the SLC18A2 gene in an individual. SLC18A2 is also known as the vesicular monoamine transporter (VMAT2). (1) is useful in studying the expression and function of SLC18A2, and in expressing the SLC18A2 protein for use in screening for candidate drugs to treat diseases related to SLC18A2 activity and in studying the effect of the variation on the biological activity of SLC18A2 as well as on the binding affinity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel genetic variants of soluble carrier family 18 (vesicular monoamine), member 2 gene useful for screening drugs to treat e.g. neuropsychiatric disorders involving monoaminergic brain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  vesicular monoamine transporter; VMAT2; polymorphic site; SNP; single nucleotide polymorphism; antiinflammatory; neuroleptic; haplotyping; genotyping; respiratory inflammatory disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human SLC18A2 allele specific oligonucleotide primer SEQ ID NO:52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   specific. The present sequence is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-SEP-2000; 2000US-232895P
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Best Local :
                                                                                                                                                                                                                                              This sequence is a PCR primer for DNA encoding the embryo-derived interleukin-related factor (EDDRF) of the invention, designated human EDIRF II. The EDIRF DNA and protein sequences (and their homologues), antibodies (Ab) specific for EDIRF, and other modulators are used:

(i) in screening and detection assays, e.g. for chromosome mapping, tissue typing or forensic studies; (ii) in diagnosis, prognosis or monitoring clinical trials; and (iii) for treating or preventing EDIRF-related diseases (especially immune, haematopoletic, differentiative, developmental or inflammatory disease, including arthritis and psoriasis. The EDIRF coding sequence, or its fragments, and the coding sequence of the se
                                                                   disease-associated mutations, also for mutagenesis), for expressing recombinant EDTRF and as source of antisense, ribozyme and peptide nucleic acids for inhibiting translation of EDIRF-derived mRNA. EDIRF is used to raise Ab (useful for detecting EDIRF, including forms with aberrant post-translational modification, for affinity purification and therapeutically) and to screen for specific modulators (e.g. peptides or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EDIRF-related disease; immune disorder; haematopoietic disorder; developmental disorder; inflammatory disease; arthritis; psoriasis; EDIRF II; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inflammatory diseases such as neuropsychiatric disorders involving monoaminergic brain systems. The present sequence represents an al specific oligonucleotide (ASO) primer for human SLC18A2, which is in the present invention.
                                                                                                                                                                                                arthritis and psuriasis. the corner (for detecting related sequences an also useful as probes and primers (for detecting related sequences an also for mutagenesis), for expressing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acid encoding embryo-derived interleukin-related factors
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В
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                                                                                                        AAA52824/c
ID AAA528
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        δ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 10
                                                     Matches
                                                                  Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                  The present sequence is one of a large number of antisense oligonucleotides designed to target different regions of the human CD44 mRNA. CD44 is a multifunctional human cell surface adhesion receptor. The oligonucleotides were analysed for effect on CD44 mRNA levels by quantitative real-time PCR analysis. Antisense oligonucleotides that inhibit CD44 expression can be used to treat CD44-associated conditions
                                                                                                                            Note: The sequence has a phosphorothioate backbone and may be either a cligodeoxynucleotide or a chimeric oligonucleotide containing 2'-methoxyethyl (2'-MOE) wings and a deoxy gap. The ISIS number given above corresponds to the oligodeoxynucleotide sequence.
                                                                                                        Sequence
                                                                                                                                                                                             including hyperproliferative disorders, such as cancer, and inflammatory conditions, such as rheumatoid arthritis. The antisense compounds hybridise to CD44 nucleic acids, thus allowing sandwich and other assays
                                                                                                                                                                                                                                                                                                                                                                       New antisense compound, that inhibits the expression of human cell surface adhesion receptor CD44, for treating hyperproliferative discrete and inflammatory conditions, such as cancer and rheumato.
                                                                                                                                                                                                                                                                                                                                      Claim 3; Page
                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-431564/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200035935-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antiinflammatory; antiarthritic; CD44 antisense inhibition; hyperproliferative disorder; cancer; inflammatory disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human CD44 antisense oligonucleotide ISIS# 18713
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-SEP-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA52824 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              rheumatoid arthritis; ss.
                                                                                                                                                                                      be easily constructed.
 16
                                                   l Similarity
11; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CD44; cell surface adhesion receptor; cytostatic; antirheumatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 CAGGTGTAGNCCA 14
|||||| || |||
3 CAGGTGCAGCCCA 15
CAGGTGTATTCCA 4
                       CAGGTGTAGNCCA 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              h 74.3%;
Similarity 84.6%;
                                                                                                        18
                                                                                                        B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cowsert LM;
                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                     76;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98US-0213719
                                                                                                        σ
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                                                                                                        P
                                                                                                                                                                                                                                                                                                                                     105pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                        5 C;
                                                              74.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18
                                                                                                        4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ВP
                                                                                                     G; 3 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                              Score 10.4;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 10.4; DB 20 Pred. No. 1.3e+04;
                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                     0 other;
                                                           .3e+04;
                                                                            DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20;
                                               Indels
                                                                        Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
                                                                           18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18;
                                                0;
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                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                               0;
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RESULT 12
AAA58857/c
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                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 11
                                                                                                                                                                                                                                                                                                                             multiwell plates; (b) a primer designed based on the chromosome marker of carey out an amplification reaction; (C) a signal corresponding to the marker is detected from the resultant camplified product to specify the discrimination Nos. of the multiwell of plates containing the clones having said marker sequence; (d) the order of the markers is changed so that the same discrimination Nos. succeed to the maximum in the specified discrimination Nos. to array the multiwell of the maximum in the specified discrimination Nos. to array the multiwell plates; (e) the clones in the multiwell plates of the specified of constituted in the specified of scrimination Nos. are mixed respectively in each wells of longitudinal confidence of the specified of constituted and the specified products; (h) the clones in the multiwell plates are cultured and the cresultant cultures are amplified by using the above primer; (g) signals are detected from the amplified products; (h) the clones in the multiwell plates are specified from the detected result; and (i) the clones are microarray is useful for gene analysis. ABL42957 to ABL45323 represent PCR primers for human chromosome 1p36-35 DNA, and ABL45323 to ABL45634 crepresent PCR primers for human chromosome 21q22.1, which are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABL43601/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 11
                                            AAA58857;
                                                                       857/c
AAA58857 standard; DNA; 19
                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                   specifically
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-MAR-2000; 2000JP-0066716
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JP2001321190-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (RIKA ) RIKAGAKU KENKYUSHO (GENO-) GENOTEX YG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            genome; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human chromosome 1p36-35 PCR primer SEQ ID NO:645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABL43601;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABL43601 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-MAR-2001; 2001JP-0068285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2002-144136/19
                                                                                                                                                        14
                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         chromosome 1p36-35; chromosome 21q22.1; genetic analysis;
                                                                                                                                                                                      CCAGGTGTAGNCC 13
                                                                                                                                                                                                                                                                                     18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Page 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     genome clones
                                                                                                                                                                                                                                                                                     BP;
                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                   claimed for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                     ω
                                                                                                                                                                                                                                                                                     Α.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  528pp; Japanese.
                                                                                                                                                                                                                                                                                   S
                                                                                                                                                                                                                                   74.3%;
84.6%;
                                                                                                                                                                                                                                                                                   Ç
                                                                                                                                                                                                                                                                                                                use in the present invention.
                                                                                                                                                                                                                                                                                   7
                                                                          ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                                                                                                                                  G
                                                                                                                                                                                                                                 Score 10.4; DB
Pred. No. 1.3e+0
                                                                                                                                                                                                                      Pred. No. 1.3
); Mismatches
                                                                                                                                                                                                                                                                                  3 T; 0 other;
                                                                                                                                                                                                                 1.3e+04;
2;
                                                                                                                                                                                                                                                  24;
                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                Length
                                                                                                                                                                                                                                                    18;
                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                   Gaps
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20-OCT-2000

(first entry)

0

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20
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AAT01071
                                                                                                                                                                                                                                                                                                                                               В
                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          oligonucleotides AAA58825-58 were used in the analysis and study of the 2g breakpoint region, in the course of the invention to identify the gene encoding a tissue repair protein. Tissue repair gene polynucleotides are useful for determining expression of mRNA in selected target tissue, e.g. for diagnosing and treating orofacial clefting. They are also useful for determining the presence
            02-MAR-1994;
                                     02-MAR-1995;
                                                              08-SEP-1995
                                                                                       W09523859-A2
                                                                                                                 Synthetic
                                                                                                                                           PCR;
                                                                                                                                                      T-cell costimulatory molecule; B7-2; T-lymphocyte; CD28; CTLA4; receptor; immunoglobulin; primer; polymerase chain reaction;
                                                                                                                                                                                         Mouse B7-1 (alternatively spliced form) PCR primer.
                                                                                                                                                                                                                        08-MAY-1996
                                                                                                                                                                                                                                                 AAT01071;
                                                                                                                                                                                                                                                                           AAT01071 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         suffering from orofacial clefting. The antibodies are also useful in the diagnosis of orofacial clefting. The polynucleotide is also useful for promoting wound healing and tissue repair.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 44; 45pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    promoting wound healing and/or tissue repair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New human and mouse nucleic acids encoding a tissue repair protein, useful for diagnosing and treating orofacial clefting, and for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Markham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-JAN-2000; 2000WO-GB00003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200040719-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tissue repair protein; orofacial cleftintissue repair; 2q breakpoint region; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oligonucleotide used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (UYLE-) UNIV LEEDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA mutations in patients suffering from, or suspected to be
                                                                                                                                           ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2000-1465983/40
                                                                                                                                                                                                                                                                                                                                               13
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                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                       CCAGGTGTAGNCC 13
                                                                                                                                                                                                                                                                                                                                              CCAGGTGGAGTCC 1
                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 4 A; 7 C; 5 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bonthron
                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99GB-0000167
            94US-0205697
                                     95WO~US02576
                                                                                                                                                                                                                                                                                                                                                                                                            74.38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            analysis and study of 2q breakpoint region
                                                                                                                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                             Pred. No. 1.30
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                            Score 10.4;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                clefting; wound healing
                                                                                                                                                                                                                                                                                                                                                                                             ..3e+04;
2;
                                                                                                                                                                                                                                                                                                                                                                                                                        DB
                                                                                                                                                                                                                                                                                                                                                                                                                        21;
                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                        Length 19;
                                                                                                                                                                                                                                                                                                                                                                                               0,
                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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RESULT 14
AAV48029
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RT-PCR was used to amplify mouse T-cell costimulatory molecule B7-1 cDNA fragments derived from murine spleen cell RNA. The primers used for primary PCR were a sense primer (AAT01068) and antisense primer (AAT01070), and for secondary PCR were a sense primer (AAT01069) and antisense primer (AAT01071). cDNA fragmer were detected that encoded a B7-1 form in which the signal peptide domain was spliced directly to the IgC-like domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAV48029 standard; DNA;
New oligo:nucleotide(s) that modulate expression of B7 proteins - used for, e.g. controlling activation and proliferation of {\tt T} cells,
                                                                                                                                                                                                                                                                                                                                       Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Murine B7-1 targetted oligonucleotide 14914.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV48029;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            i.e. the IgV-like domain was deleted (see AAR82900). Another cDNA fragment coded for a B7-1 form in which the IgV-like domain was spliced directly to the transmembrane domain, i.e. the IgC-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel T cell co-stimulatory molecules - corresponding to naturally occurring alternatively spliced forms of T cells co-stimulatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Borriello
                                                     WPI; 1998-387783/33
                                                                                                                                                             31-DEC-1996;
                                                                                                                                                                                                                               09-JUL-1998
                                                                                                                                                                                                                                                                  WO9829124-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                cell proliferation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1995-320574/41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BGHM ) BRIGHAM & WOMENS HOSPITAL (DAND ) DANA FARBER CANCER INST.
                                                                                                                          (ISIS-) ISIS
                                                                                                                                                                                              16-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 domain was deleted (see AAR82902).
                                                                                                                                                                                                                                                                                                                                                                                           lomo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 CCAGGTGTAGNCC 13
                                                                                        CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CCAGGTGAAGTCC 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 BP; 5 A; 6 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 or variants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   B7; T cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 43; 111pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                          PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Freeman
                                                                                                                                                             96US-0777266
                                                                                                                                                                                            97WO-US23270
                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                  /*tag= a
/note= "Phosphorothioate linkages'
                                                                                                                                                                                                                                                                                                                        /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        74.3%;
84.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   inflammation; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ВP
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Pred. No. 1.3e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cell activation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
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particularly for treatment, diagnosis and prevention of inflammation

English

Example 1; Page 36;

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RESULT 15
AAA91878/c
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Best Local
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                                                The present invention relates the diagnosis of genetic susceptibility for end-stage renal disease (ESRD). The method involves analysing a polynucleotide sample for a single nucleotide polymorphism (SNP) associated with an altered susceptibility for ESRD. The method allows early detection of ESRD and hence effective delay or ideally, prevention of ESRD is made possible. The present sequence is a primer used to amplify bases 66 to 265 of human TGF bl gene. Polymorphisms the TGF bl gene are known to be a probable trigger for renal apoptosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     particularly useful for treatment and prevention of inflammation and autoimmune diseases, e.g. asthma, (juvenile) diabetes, myasthenia gravis, Grave's disease, rheumatoid arthritis, allograft rejection, psorilasis, (systemic) lupus erythematosus, multiple sclerosis, contact dermatitis, rhinitis, allergy, cancer and metastases. The oligonucleotides may also be used to manipulate T cell activation ex vivo; to determine or detect B7 protein expression; for diagnosis; as assay and purification reagents, and to study physiological roles of B7 proteins.
Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                    Diagnosing genetic susceptibility for end-stage renal disease using single nucleotide polymorphisms, involves analyzing sample obtained from subject to detect genetic polymorphism in the sample polynucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antisense primer to amplify bases 66 to 265 of human TGF b1 gene.
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                                                                                                                                                                                                                                                                                                                          Example 3; Page 36; 73pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Moskowitz
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (DZGE-) DZGENES LLC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24 - AUG - 2000
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BP; 6 A; 5 C; 6 G; 3 T; 0 other;
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Pred. No. 1.3e+04;
Pred. No. 1.3e+04;
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Search completed: June 21, 2003, Job time: 78.7812 secs
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84.6%;
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and is derived
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       No. is the number of results predicted by chance to have a greater than or equal to the score of the result being printed, derived by analysis of the total score distribution.
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/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
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/cgn2_6/ptodata/1/ina/FB_COMB.seq:*
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US-08-585-684B-820
US-08-855-684B-821
US-09-038-073-821
US-09-033-073-821
US-09-033-77-266A-68
US-08-77-266A-68
US-08-702-525-61
US-09-907-843-48
PCT-USS-0257-61
US-07-998-289B-26
US-08-859-998-1077
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US-09-139-617-15
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   -495-695B-25
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                                 Sequence 48, Appl Sequence 820, App Sequence 821, Appl Sequence 14, Appl Sequence 61, Appl Sequence 1077, Appl Sequence 1077, Appli Sequence 1077, Appli Sequence 1077, Appli Sequence 1077, Appli Sequence 15, Appli Sequence 58, Appli Sequence 58,
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1225 225 138 137 17 17 17 17 17 17 17 17 17 17 17 17 17	19, Appl	Appli	, Appl	, Appl	Appl	. Appl	20, Appl	19, Appl		`	543, App		17, Appl	139, App	•	•	~	

## ALIGNMENTS

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EARLIER APPLICATION NUMBER: 60/047,352
EARLIER FILING DATE: 1997-05-21
NUMBER OF SEQ ID NOS: 871
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 48
LENGTH: 15
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Gene Expression Profiles TITLE OF INVENTION: Cancer Cells FILE REFERENCE: 01107.74664
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APPLICANT: Vogelstein, Bert
APPLICANT: Zhang, Lin
APPLICANT: Zhou, Wei
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                                                                                          TITLE OF INVENTION: ITILE OF INVENTION: ITILE OF INVENTION: INUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo sapiens
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
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o. 6333152
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                                                                                                                                                             McSwiggen, James
                                                                                                                                                                                 Jarvis, Thale
                                                                                                                                                                                                  Stinchcomb, Daniel T.
                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                          METHOD AND REAGENT FOR THE INDUCTION OF GRAFT TOLERANCE AND REVERSAL OF IMMUNE RESPONSES: 2751
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91.7%;
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Pred. No.
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                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                          SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/585,684B
                              FILING DATE: January 16, 1996 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Stinchcomb, Danie APPLICANT: Jarvis, Thale APPLICANT: McSwiggen, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/00
FILING DATE: July 7, 1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600.
                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                         STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (213) 955-0440
TELEX: .67-3510
APPLICATION NUMBER: 6
FILING DATE: July 7,
                                                                                                                                                                                                                           CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P O
                                                                                                             OPERATING SYSTEM:
                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
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11; Conserv
                                                                                                                                                                                               90071
                                                                                                                                                                                                                                                        E: Lyon & Lyon
633 West Fifth Street
Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Application US/08585684B
                                                                                                                                                                                                          U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                       Stinchcomb, Daniel T.
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                                                                                                                          IBM Compatible
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                                                                                                                                                                                                                                                                                                                         METHOD AND REAGENT FOR THE INDUCTION OF GRAFT TOLERANCE AND REVERSAL OF IMMUNE RESPONSES 2751
                                                                                                             IBM P.C. DOS 5.0
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               60/000,951
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Pred. No 1
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Query Match

74.38;

Score 10.4;

DB 4;

Length 15;

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RESULT 4
US-09-038-073-820/c
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US-09-038-073-820
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 821:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                             TELEFAX: (213) 955-044
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                               REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                 OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: FASTESQ Version 1.5 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/038,073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF
                                                                                                                                                                                                                                                   FILING DATE:
PRIOR APPLICATION DATA:
                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, Jar
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            STRANDEDNESS:
TOPOLOGY: li
                                                                                                                                                                            NAME: Warburg, Richard REGISTRATION NUMBER: 3
                                                                                                                                                                                                                       FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
                                                       LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          820
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                                      15 base pairs
ucleic acid
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                                                                                                                  (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Stinchcomb, Daniel T.
                                                                                                                                                                                                                                                                                                                                         IBM Compatible
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             linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            74.3%;
84.6%;
                                                                                                                                                                            32,327
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                                                                                                                                                              218/078
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Pred. No. 1.5e+03
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RESULT 6
US-09-213-719-14/c
Sequence 14, Application US/09213719B;
Patent No. 6150162;
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                 GENERAL INFORMATION:
APPLICANT: C. Frank Bennett
APPLICANT: Lex M. Cowsert
                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
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Matches 11; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 821,
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TITLE OF INVENTION: ANTISENSE MODULATION OF CD44 EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NEORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER RÉADABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ Version 1.5 CURRENT APPLICATION DATA:
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 633 West Fif
STREET: Suite 4700
CITY: Los Angeles
STATE: California
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                                                                                                                                                                                                                                                                                                                                                           Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Warburg, Richard REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
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                                                                                                                                                                                                                                      CCAGGTGAAGTCC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          67-3510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
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                                                                                                                                                                                                                                                                                                                                                         74.3%;
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ER: 218/078
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                                                                                                                                                                                                                                                                                                                         Score 10.4; UD ,
Pred. No. 1.5e+03;
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                                                                                                                                                                                                                                                                                                                                                                       Length 15;
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                                                                                       Query Match
Best Local Similarity
Matches 11; Conser
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (609) 779-84 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                           NAME: Jane NUMBER: 32,257
REGISTON NUMBER: 15'
REFERENCE/DOCKET NUMBER: 15'
TELECOMMUNICATION INFORMATION:
(609) 779-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: Decembe CLASSIFICATION: 536 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Clarence Frank Bennett APPLICANT: Timothy A. Vickers TITLE OF INVENTION: Oligonuclectic TITLE OF INVENTION: Methods for the control of the contro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Cherry Hill
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
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>. 6077833
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2 CCAGGTGAAGTCC 14
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                                                                                         Conservative
                                                                                                                                                                                                              Yes
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- NO: 68:
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84.6%;
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84.6%;
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; OTHER INFORMATION: Antisense Oligonucleotide US-09-213-719-14
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CURRENT APPLICATION NUMBER: US/09/213,719B
CURRENT FILING DATE: 1998-12-17
NUMBER OF SEQ ID NOS: 91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Law Offices of Jane Massey Licata STREET: 210 Lake Drive East, Suite 201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oligonucleotide Compositions and Methods for the Modulation of the
                                                                                                                                                                                                                                                                                                                                                                  US/08/777,266A
                                                                                                                                                                                                                                 ISPH-0201
                 Score 10.4;
Pred. No. 1
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Pred. No. 1.5e+03;
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   Mismatches
                 1.5e+03
                                DB
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RESULT 9
US-08-702-525-61
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US-08-205-697A-61
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                                                                                                                                                                                                                                                                                                                                                                                      Matches
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/205,697A
FILING DATE: 02-Mar-1994
ATTORNEY/AGENT INFORMATION:
                                                                      NUMBER OF SEQUENCES: 65
CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD
                                                                                                                       TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: BW TELECOMMUNICATION INFORMATION: TELEPHONE: (617)227-7400
                                                                                                                                                       APPLICANT:
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MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS:
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               COUNTRY:
                               STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TORNEE / AGENT AND E. NAME: Mandragouras, Amy E. REGISTRATION NUMBER: 36,207
REGISTRATION NUMBER: BWI-120
                                                           STREET:
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02109-1875
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                                                         28 State Street
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                                                                                                                                                                                 Sharpe, Sharpe
Borriello, Francescopaolo
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                                                                                                           Nadler, Lee
PARTION: NO. 6294660el Forms of T Cell Costimulatory
VENTION: Molecules and Uses Therefor
                                                                                                                                                                   Freeman,
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84.6%;
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                                                                                                                                                                 Gordon
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Pred. No. 1
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US-09-907-843-48/c
US-09-907-843-48/c
: Sequence 48, Application US/09907843
: Patent No. 6440739
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US-09-326-186B-68
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APPLICANT: Bennett, Clarence Frank
APPLICANT: Vickers, Timothy A.
                                                                                                                                                                              Matches
                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 68
LENGTH: 20
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Best Local :
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PRIOR FILING DATE: 1996-12-31
NUMBER OF SEQ ID NOS: 226
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: ISPH-0376
CURRENT APPLICATION NUMBER: US/09/326,186B
CURRENT FILING DATE: 1999-06-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Vickers, Timothy A.

TITLE OF INVENTION: Oligonucleotide Compositions and Methods for the
TITLE OF INVENTION: Modulation of the Expression of B7 Protein
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                           TYPE: DNA
                                                                                                                                                                                                                                                                                                                                          LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: BW
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 0 FILING DATE: 02-mar-1994 ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Floppy disk
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                                                                                                                          1 CCAGGTGTAGNCC 13
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l Similarity 84.6%;
ll; Conservative
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                                                                                                       CCAGGTGAAGTCC 14
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                                                                                                                                                                              Conservative
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SYSTEM: PC-DOS/MS-DOS
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84.68;
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Pred. No. 1.5e+03;
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                                                                                                                                                                                                .5e+03;
                                                                                                                                                                                                               DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 4;
                                                                                                                                                                                                            Length 20;
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                                                                                                                                                                              Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCT-US95-02576-61 Sequence 61, Application PC/TUS9502576
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-907-843-48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 48
LENGTH: 20
TYPE: DNA
                                                   Matches
                                                                                 Query Match
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                                                                                                                                                                                                        TELEPHONE: (617)227-7400
TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 61
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/907,843
CURRENT FILING DATE: 2001-07-17
NUMBER OF SEQ ID NOS: 87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: 'SUSAN M. Freier
TITLE OF INVENTION: ANTISENSE MODULATION OF GLIOMA-ASSOCIATED ONCOGENE-2 EXPRESSION
FILE REFERENCE: RTS-0279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Antisense Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                               COMPULE.

OPERATING SYSTEM: PC-DUS/No LUC
SOFTWARE: ASCII Text
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:
                                                                                                                               STRANDEDNESS:
TOPOLOGY: lin
MOLECULE TYPE:
                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: BW
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 0 FILING DATE: 02-Mar-1994 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 65
CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                  / Match 74.3%;
Local Similarity 84.6%;
nes 11; Conservative
                                                                                                                                                                             TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                            NAME: Mandragouras, Amy E. REGISTRATION NUMBER: 36,20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
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                  1 CCAGGTGTAGNCC 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    60 State Street, suite 510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Frank Bennett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IBM PC compatible
                                                                                                                                                 linear
                                                                                                                                oligonucleotide
                                                                                                                                                             single
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                                                   0,
                                                Score 10.4; DB 5;
Pred. No. 1.5e+03;
0; Mismatches 2;
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Pred. No. 1.
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                                                                                                                                                                                            RESULT 14
                                                                         Sequence 1077, Application US/08859998
Patent No. 5994076
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Jokhadze, George
APPLICANT: Bibilashvilli, Robert
                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 23 base pairs
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                               TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                   IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION: TELEPHONE: 212-527-7700
TELEFAX: 212-753-6237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: METITLE OF INVENTION: RENUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                    CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Robinson, Joseph R
REGISTRATION NUMBER: 33,448
REFERENCE/DOCKET NUMBER: 06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/07/998,289B FILING DATE: 30-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                         CLONE:
                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM:
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   ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            805 Third Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Black, Bruce C
Taylor, Martin
Heckel, David G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              S
                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                       Heliothis virescens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
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Fish & Richardson, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  single
                                                                                                                                                                                                                                                                                                                        74.3%;
                                 EXPRESSION 1375
                                                               METHOD OF ASSAYING DIFFERENTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Resistance
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                                                                                                                                                                                                                                                                                                                        Score 10.4;
Pred. No. 1
                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                                                                                                                                                                                                                                                                                           DB
                                                                                                                                                                                                                                                                                                                                       Length 23;
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Gaps

COMPUTER READABLE FORM: MEDIUM TYPE: Diskette

COUNTRY: UZIP: 94025

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STREET: 2200 Sand CITY: Menlo Park

2200 Sand Hill Road, Suite 100

CA

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; OTHER INFORMATION: oligonucleotide primer
US-08-859-998-1077
                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 15
US-08-765-332-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 11
                                                                                                                                                                                                                                                                                                                                                                                       Patent No. 6025132
GENERAL INFORMATION:
APPLICANT: JANNES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY_AGENT INFORMATION:

NAME: Field, Bret E.

REGISTRATION NUMBER: 37,620

REFERENCE_POCKET NUMBER: 0909

TELECOMMUNICATION INFORMATION:

TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 415-854-0875
INFORMATION FOR SEQ ID NO:
                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compartible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS
                               APPLICATION NUMBER: US/08/765,332
FILING DATE: 23-DEC-1996
                                                                                                                                                                       COUNTRY: UZIP: 22201
                                                                                                                                                                                                       STATE: D.C
                                                                                                                                                                                                                                        STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
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                                                                                                                                                                                                                      CITY: WASHINGTON
                                                                                                                                                                                                                                                         ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FH: 24 base pairs nucleic acid NDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 CCAGGTGTAGNCC 13
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                                                                                                                                                                                                                                    E: NIXON & VANDERHYE.P.C.
1100 NORTH GLEBE ROAD, 8TH FLOOR
                                                                                                                                                                                                                                                                                                                                                      ROSSAU, RUDI
VAN HEUVERSWYN, HUGO
                                                                                                                                                                                       USA
                                                                                                                                                                                                                                                                                                                                                                                     JANNES, GEERT
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                                                                                                                                                                                                                                                                                                     SIMULTANEOUS DETECTION, IDENTIFICATION AND DIFFERENTIATION OF EUBACTERIAL TAXA USING A HYBRIDIZATION ASSAY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Windows95
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Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 2; Length 24;
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В
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Best Local Similarity
""tches 11; Conserva
Search completed: June 22, 2003, 00:01:36 Job time: 18.625 secs
                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: EP 948
FILING DATE: 24-JUN-1994
ATTORNEY/AGENT INFORMATION:
NAME: SADOFF, B.J.
REGISTRATION NUMBER: 36,66
REFERENCE/DOCKET NUMBER: 1
                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                          MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                             ANTI-SENSE:
                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: 703-816-4091
                                                                                                                                                                                                                                                   LENGTH: 24 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                    TELEPHONE: 705 CT TO NO:
                                                                                                                                                                                                                                        TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: WO PCT/EP95/02452
                                                          9 CCAGGTGTTGTCC 21
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                                                                                                                    Conservative
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24-JUN-1994
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                                                                                                                                   74.3%;
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                                                                                                                                                                                                                                                                                                                     4:
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                                                                                                                   Score 10.4; DB 3; Pred. No. 1.5e+03; 0; Mismatches 2;
                                                                                                                                                 Length 24;
                                                                                                                    Indels
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Gaps

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AZ346490 1M0081004 AZ832832 2M0113D06

R63848 yi21a09 BE739587 601556558

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Post-processing: Minimum Match
| Maximum Match
Listing first
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Maximum DB
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Sequence:
                                                                          Result
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                                                                                                        is the number of results predicted by chance to have a ater than or equal to the score of the result being printed, rived by analysis of the total score distribution.
                                                                 Match
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                                                                          Query
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                                                                                                                            AUTHORS
                                                                AUTHORS
                                                                                                                                                                                                                                                                                                               Strizhov,N., Li,Y., Rosso,M. and Weisshaar,B. Direct Submission Submitted (17-JUN-2002) Weisshaar B., Max-Planck-Institut
                                            Rosso,M., Strizhov,N., Li,Y., Reiss,B., Dekker,K. and Weisshaar,B. A new Arabidopsis thaliana T-DNA mutagenised population (GABI-Kat) for flanking sequence tag based reverse genetics
                                                                                     and Weisshaar,B.

A pipeline for automated high-throughput generation of FSTs (flanking sequence tags) from Arabidopsis thaliana T-DNA transformed lines
                                                                                                                                                 Arabidopsis thaliana
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicottyledons; core eudicotts;
Spermatophyta; Magnoliophyta; Processing Corp. Arabidocaid
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thale cress.
                              Unpublished 3 (bases 1
                                                                                                                                                                                                  genomic survey sequence.
AL766269
AL766269.1 GI:21519402
                                                                                                                                                                                                                             AL766269 41 bp DNA linear GSS 18-JUN-2002 Arabidopsis thaliana T-DNA flanking sequence GK-209A10-014557,
                                                                                Unpublished
                                                                                                                           Strizhov, N., Li, Y., Rosso, M., Viehoever, P.,
                                                                                                                                             Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis
                           (bases 1 to 41)
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AZ872071
AZ628215
AI719563
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AZ445734
AA127131
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C21079 HUNGS000259
T93720 ye09g01.51
AU105239 AU105239
AU107231 AU107231
BM182676 fv60f03 y
N34794 yy16g04.51
AZ864400 2M0173824
AZ772618 1N0583H09
AZ782618 1N0583H09
AZ588484 1M0390M07
AL768826 Arabidops
AZ78618160 1M0369005
AZ461229 1M0367904
AZ601449 1M0419A16
AZ875800 2M019G04
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AU104849 AU104849
BH233070 100617100
AI1219622 9910007 x
AZ491277 IM0324E14
AL464465 T bircei
AZ924604 4906 IC29
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AA910920 ok87h09.s
AZ872071 ZM0185E13
AZ872071 ZM0185E13
AZ628215 IM0440007
AI719563 as54409.x
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AA510334 vg32c11 r
BIZ59862 602971524
BH791916 SALK_0619
AZ579484 IM0367806
AZ445734 IM0241M16
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T68434 yc42d01, s1

AZ492224 IM0326B13

Saedler, H.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   German Plant Genomics program designated 'GABI'. Information on line availability can be found at: http://www.mpiz-koeln.mpg.de/GABI-Kat/. Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Zuechtungsforschung, Carl-von-Linne-Weg 10, Koeln, 50829, Germany This sequence is recovered from the left border of the T-DNA. It indicates an insertion within the locus defined by clone f26h11. The sequences are generated at the MPI for Plant Breeding Research in the context of the GABI-Kat project. GABI-Kat is part of the
                                                                                                                             Plate: 0151 row: P column: 19
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Rei,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AZ850196 37 bp DNA IIIIEGI WSS 2. 2M0151P19R Mouse 10kb plasmid UUGC1M library Mus musculus clone UUGC2M0151P19 R, DNA sequence.
                                                              High quality sequence stop:
Location/Qualifiers
                                                                                                                                                       Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
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Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                      Rm.
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                                                                                                            Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                            Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                     plasmid inserts
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus
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/note="PCR was performed on DNA from Arabidopsis thaliana
plants (T1) which were transformed with the T-DNA from
vector pACI61. The lines contain one or more T-DNA
insertions. The DNA fragment(s) resulting from the PCR
were directly sequenced to determine the genomic sequence
flanking the insertion. Sequences displaying significant
similarity to the A. thalaina nuclear genome sequence were
processed for submission. T-DNA derived sequences were
removed"
                                                                                                                                                                                                                                                                                                 Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:3702"
/clone="GK-209A10-014557"
/organism="Mus musculus"
/strain="C57BL/6J"
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/strain="Columbia 0"
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91.7%;
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Pred. No. 3.1e+04;
                                                                                                                                                                                Std Error: 0.00
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PO Box 6104, Newark, PO Box 6104, Newark, PO Box 631 2638
Tel: 302 631 2607
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G-6c8.f Maize Random Small-insert clone G-6c8 both, DNA sequence.
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Zea mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Meyers, B.C., Tingey, S.v. and Acceptable Abundance, distribution and transcriptional
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BH129921
BH129921.1
                                                                                                                                                                                              Email: Michele.morgante@usa.dupont.com
Sequences were trimmed to include only high quality bases;
and reverse reads were assembled when significant overlaps
                                                                                                                                                                                                                                                                                                                                                                       Suite 200
                                                                                                                                                                                                                                                                                                                                                                                          Contact: Morgante
                                                                                                                                                                                                                                                                                                                                                                                                                21475670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACC
                                                                                                                                                     Seq primer: Ml3univ
                                                                                                                                                                                  detected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    clade; Panicoideae; Andropogoneae;
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                                                                                                                                    shotgun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance. "  7 \text{ c} 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil473214) gb|AFT29072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and mith adaptors complementary to the insert adaptors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA
                                    /organism="Zea mays"
/strain="B73"
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
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/note="Vector: PWD42nv; Purified genomic DNA from N
                   /db_xref="taxon:4577"
                                                                                                          Location/Qualifiers
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Plate: LLAM8772 row: p column: 23
High quality sequence stop: 38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Jeffrey Green M.D.
cDNA Library Preparation: Life Technologies,
cDNA Library Arrayed by: The I.M.A.G.E. Conso
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
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601232196F1 NCI_CGAP_Mam6
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National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                     providing samples: Jeffrey Green,
                                                                                                                                                                     /note="Organ: mammary: Vector: pCMV-SPORT6; Site_1: SalI; Site_2: NotI; Cloned unidirectionally: Primer: Oligo dT. Library constructed by Life Technologies. Investigator providing samples: Jeffrey Green, M.D., NIH"
                                                                                                                                                                                                                                           /clone="IMAGE:3596038"
/clone_lib="NCI_CGAP_Mam6"
/sex="female, virgin"
/tissue_type="infiltrating ductal carcinoma'
/dev_stage="5 months"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                    /strain="FVB/N"
/db_xref="taxon:10090"
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/dev_stage="seedling"
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Gilbert Smith, Ph.D.
Tissue Procurement: Gilbert Smith, Ph.D.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be.
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Hordeum vulgare subsp. vulgare
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Pooldea;
Triticeae; Hordeum.
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vulgare leaves vegetative stage Hordeum vulgare subsp. vulgare
clone baak12m18, mRNA sequence.
                                                                                                                                 NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae. 1 (bases 1 to 48)
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Research Institute for Bioresources
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                                                                                                                   Contact: Robert Strausberg, Ph.D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               URL:http://www.rib.okayama-u.ac.jp/barley/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: kazsato@rib.okayama-u.ac.jp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Hordeum vulgare subsp. vulgare"
/cultivar="Akashinriki"
/db_xref="taxon:112509"
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BE739587
BE739587.1 GI:10153579
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National Institutes of Health, M
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Plate: LLAM11038 row: p column: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Plate: LLCM493 row: d column:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (1999)
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http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CDNA Library Preparation: CLONETECH Laboratories, Inc. CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          numan
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/clone_lib=*NIH_MGC_58"
/clone_lib=*NIH_MGC_58"
/tissue_type="hypernephroma"
/lab_host="PH10B (T1 phage-resistant)"
/note=*Organ: kidney; Vector: pDNR-LIB (Clontech); Site_1:
/note=*Organ: kidne
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Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Library constructed by Life Technologies. Investigator
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                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/db_xref="taxon:9606"
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/dev_stage="5 months"
/lab_host="DH10B"
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Pred. No. 7.
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y121a09.rl Soares placenta Nb2HP Homo sapiens cDNA clone
IMAGE:139864 5' similar to SP:TSC2_MOUSE Q00992 PUTATIVE REGULATORY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information
Trace considered overall poor quality
Possible reversed clone: similarity on wrong strand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 (bases 1 to 28)
Hillier, L., Clark, N.,
M., Hultman, M., Kuca
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Fax: 314 286 1810
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (1995)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seq primer: M13RP1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: est@watson.wustl.edu
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                                                                                                                                   constructed
                                           and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization. Library
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                                                                                   double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the
                                                                                                                                                                                                                                                                                                                      /db_xref="GDB:550462"
/db_xref="taxon:9606"
/clone="IMAGE:139864"
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                                                                                                                                                                                                                                                                             /sex="Female"
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Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,
ing,T., Soares,M., Tan,F., Trevaskis,E., Waterst
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by Bento Soares and M.Fatima Bonaldo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0081 row: 0 column: 04
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AZ346490 31 bp DNA linear GSS 29-SEP-200 1M0081004R Mouse 10kb plasmid UUGC1M library Mus musculus genomic UUGC1M0081004 R, DNA sequence.
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sclurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus
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AZ346490.1 GI:10425727
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
                                                                                                                             (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNDA2 (gil4732114[gb]AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    inserts
with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                      /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Mus musculus"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                     Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone_lib="Mouse 10kb plasmid UUGClM library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone="UUGC1M0081004"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            'sex="Male"
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RESULT 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0113 row: D column: 06
Seq primer: CACACAGGAAACAGCTATGACC
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Dunn, D., Aoyagi, A.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert B.
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was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwp42 (gi147321141gb1AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated
                                                                                                                                                                                                                                                                                                (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA
                                                                                                                                                                                                                                                                                                                                                                                     /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Mus musculus"
/strain="C57BL/GJ"
/db_xref="taxon:10090"
/clone="UUGC2M0113D06"
/clone_lib="Mouse 10kb plasmid UUGCIM library"
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                                                                                  with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was anneale
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chordata; Craniata; Vertebrata; Euteleostomi;
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Pred. No. 1.1e+05;
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RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Trace considered overall poor quality
Possible reversed clone: similarity on wrong strand seq primer: -40ml3 fwd. Ef from Amersham
High quality sequence stop: 1.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost,S., Krizman,D., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Mart,J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Theising,B.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AA677571

46 bp mRNA linear EST 19-DEC-1997
zj66f03.sl Soares_fetal_liver_spleen_lNFLS_S1 Homo sapiens cDNA
clone_IMAGE:455261 3' similar to TR:G535185 G535185 FIBRINOGEN-LIKE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 CCAGGTGTAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   White,Y., Wylie,T., WaterstowashU-NCI human EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PROTEIN. ;, mRNA sequence. AA677571
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
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                                                                                         Similarity
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314 286 1810
                                                                                                                                                              16
                                                                       Conservative
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                                                                                                                                                                                            double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Pac I and cloned into the Pac I and Eco RI sites of the modified pTTT3 vector. Library went through one round of normalization. Library
                                                                                                                                                                                                                                                                                                          /dev_stage="20 week-post conception fetus"
/lab_host="PH10B (ampicillin resistant)"
/note="Organ: Liver and Spleen: Vector: pT773D (Pharmacia)
with a modified polylinker; Site_1: Pac I; Site_2: Eco RI;
This is a subtracted version of the original Soares fetal
                                                                                                                                                        constructed by Bento Soares and M.Fatima Bonaldo." 6 c 9 g 15 t
                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                          /sex=
                                                                                                                                                                                                                                                                                                                                                                                                                      /clone_lib="Soares_fetal_liver_spleen_1NFLS_S1"
                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone="IMAGE:455261"
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100.0%;
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                                                                                        Score 10;
Pred. No.
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Pred. No.
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                                                                                                        Length 46;
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                                                                      Gaps
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RESULT 13
BI259862/c
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DEFINITION

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BI259862 31 bp \mathfrak{m} 602971524F1 NIH_MGC_12 Homo sapiens
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Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Trace considered overall poor quality
Possible reversed clone: similarity on wrong strand
Seq primer: -28ml3 rev2 ET from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: 314 286 1800 Fax: 314 286 1810
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Marra,M., Hillier,L., Allen,M., Bowles,M., Dietrich,N., Dul
Geisel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., Morris,M.
Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AA510334 28 bp mRNA linear EST 08-JUL-1997 vg32c11.r1 Soares_mammary_gland_NbMMG Mus musculus cDNA clone IMAGE:863060 5' similar to SW:OST4_CHICK P48440 OLIGOSACCHARYL TRANSFERASE 48 KD SUBUNIT ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               High quality sequence stop: 1.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus
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Bonaldo."
a 9 c
                                                                                                                                                                                                                                                                                                                                                                                                                                           the Not I and Eco RI sites of the modified pT7T RNA provided by Dr. Minoru Ko, Wayne State Univ. constructed and normalized by Bento Soares and M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              adaptors (Pharmacia),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RI; 1st strand cDNA was primed with a primer [5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note="Organ: mammary gland; Vector: pT7T3D-Pac (Pharmacia /note="Organ: mammary gland; Vector: pT7T3D-Pac (Pharmacia ) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; 1st strand cDNA was primed with a Not I - oligo(dT)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /tissue_type="mammary gland"
/dev_stage="4 weeks"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="IMAGE:863060"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone_lib="Soares_mammary_gland_NbMMG"
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                                                                                                                                                                                                                                                                                         70.0%;
78.6%;
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                                                                                                                                                                                                                                                                0;
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                             mRNA
nRNA linear EST 17-JUL-2001 CDNA clone IMAGE:5111260 5',
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                                                                                                                                                                                                                                                                                                              Length 28;
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RESULT 14
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                                                                                                                                 1 (bases 1 to 42)
Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R.,C.J., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L.,Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in the
                                                                                                                                                                                                                                                                                                                                                                                                  BH791916
SALK_061994.25.20.x Arabidopsis thaliana TDNA insertion Arabidopsis thaliana genomic clone SALK_061994.25.20.x,
                Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGNAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
                                                                                                  Unpublished (2001)
                                                                                                                                                                                                                                      Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                              BH7,91916.1
GSS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                      Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                       thale cress.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             http://image.llnl.gov
Plate: LLAM11269 row: o column:
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National Institutes of Health, Mammalian
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (1999)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Torrey Pines Road, 453 4100 x1752
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/lab_host="DH10B"
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/clone_lib="NIH_MGC_12"
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/db_xref="taxon:9606"
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Insert Length: 10000 Std Erro
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University of Utah
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                                                                                                                                                                                                                                                                                                                                                                             Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                and Wright, D., Weiss, R.
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/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
                 /clone_lib="Mouse 10kb plasmid UUGC1M library"
/sex="Male"
                                                   /db_xref="taxon:10090"
/clone="UUGC1M0367B06"
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/strain="Columbia 0"
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 /lab_host="E.
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S., Mahmoud, M., Meenen, E.,
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AR006801
AR006801.1
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12; Conserv
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12; Conserv
                                                                                           1 (bases 1 to 17)
Newman, R.A., Hanna, N. and Raab, R.W.
Recombinant antibodies for human therapy
Patent: US 5750105-A 51 12-MAY-1998;
                                                                                                                                Unclassified
                                                                                                                                                   Unknown.
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                                                                                                                                                                                                                                                                                                                                                                                          Probes and decoder oligonucleotides patent: WO 0216649-A 4261 28-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                 Gunderson, K.
                                                                                                                                                                                                                                                                                                                                                                                                                                           synthetic
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                                                                                                                                                                                                                                                                                                                                                                                     Illumina,
                                                                                                                                                                                                                                                                                                                                                                                                                                artificial sequences
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                                                     /organism="unknown"
5 c 6 g
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/db_xref="taxon:32630"
/note="Computer Generated Probe
5 c 8 g 6 t
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                 81.4%;
85.7%;
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92.3%;
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Pred. No. 2.5e
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Pred. No.
        Score 11.4; DB 6;
Pred. No. 5.8e+04;
0; Mismatches 2;
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161180
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1 CAAGGTCCAGNCCA 14
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Hanna,N., Newman,R.Anthony. and Reff,M.Elliot.
Recombinant anti-CD4 antibodies for human ther
Patent: US 6136310-A 44 24-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                 AR135409
Sequence 44 from patent US
AR135409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 (bases 1 to 17)
Newman,R.A., Hanna,N. and Raab,R.W.
Recombinant antibodies for human therapy
Patent: US 5756096-A 1 26-MAY-1998;
Location/Qualifiers
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Unclassified.
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Pred. No. 5.8e+04;
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Pred. No. 5.8e+04;
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REFERENCE
AUTHORS
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178729/c
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                                                                                                                                                                                                                                                                           12;
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Newman,R.A., Hanna,N. and Raab,R.W.
Recombinant antibodies for human therapy
Patent: US 5681722-A 51 28-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Newman, R.A., Hanna, N. and Raab, R.W. Recombinant antibodies for human therapy Patent: US 5658570-A 5 19-AUG-1997;
  1 CAAGGTCCAGNCCA 14
                     12; Conservative
                                                                                                            1 (bases 1 to 17)
Newman,R.A., Hanna,N. and Raab,R.W.
Recombinant antibodies for human therapy
Patent: US 5693780-A 5 02-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unknown
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Pred. No. 5.8e+04;
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AR151630
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AR103422
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AR151630
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AR103422
AR103422.1
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Sequence 16
AX250348
AX250348.1
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1 (bases 1 to 22)
Conklin,D.C., Foster,D.C. and Gao,Z.
Kunitz domain polypeptide and materials and methods for making
Patent: US 6087473-A 13 11-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unclass...
( bases 1 to 18)
Reinl,S.J. and Turpen,T.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent: WO 0168682-A 16 20-SEP-2001;
Large Scale Biology Corporation (US)
13 from patent US
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13 from patent US 6087473.
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1. .22
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7 c 7 g
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/note="primer"
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Pred. No. 5.7e+04;
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Conklin,D.C., Foster,D.C. and Gao,Z.
Kunitz domain polypeptide and materials
Patent: US 6232098-A 13 15-MAY-2001;
Location/Qualifiers
                                                   1 (bases 1 to 26)
Newman, R.A., Hanna, N. and Raab, R.W.
Recombinant antibodies for human therapy
Patent: US 5750105-A 20 12-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                 Boehm, T. and Dear, N.T.
Calpain protease 12
Patent: WO 02027/75-A 19 10-JAN-2002;
BASF AKTIENGESELLSCHAFT (DE)
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AR135378
           1 CAAGGTCCAGNCCA 14
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Newman,R.A., Hanna,N. and Raab,R.W.
Recombinant antibodies for human therapy
Patent: US 5681722-A 20 28-OCT-1997;
                                                                                                                                                                           Unknown
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                                                                                                                                                                  Unclassified
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Hanna,N., Newman,R.Anthony. and Reff,M.Elliot.
Recombinant anti-CD4 antibodies for human ther
Patent: US 6136310-A 13 24-OCT-2000;
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11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4	11.4
81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4.	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4	81.4
34	34	32	30	30	30	30	29	27	27	27	27	26	26	26	26	26	26	26	25	24	22	22	22	22	20	20	20	18	<u>1</u> 8	18	17	17	17	17	17
21	19	17	19	17	15	15	22	21	19	18	16	19	19	19	18.	18	18	14	18	24	24	22	21	21	22	22	20	22	22	18	19	19	19	18	18
AAA78273	AAV61365	AAT18053	AAV33320	AAT16179	AAQ56586	AAQ67195	AAH47905	AAA09964	AAV52801	AAT92804	AAQ87238	AAV05652	AAV23762	AAV31384	AAT62871	AAT92201	AAT95124	AAQ35905	AAT75424	ABA99782	ABL55000	AAD14335	AAZ46336	AAA50117	ААН99913	AAS12663	AAV64442	AAF24764	AAS45544	AAT43427	AAV05683	AAV23793	AAV31430	29	AAT91566
Human Ig H chain s	Human immunoglobul	prin		Humanised antibody	μ.	Humanized 1308F VH	IgE antibody L cha	for		HS1 for	Primer for leader	onkey .Vh1	r Anti-		Human or monkey VH	QJ.	Human or monkey Ig	Human/monkey heavy	Primer for human a	e capn5 se	lympho	Human zkun5 DNA am	zkun5 F	erine	r used to am	BAX	Mouse ada gene rev	5' PCR primer.used	Tumour-specific Ig	VH1 family specifi	Human/monkey heavy		(n	$\circ$	Cynomolgus monkey

## ALIGNMENTS

RESULT 1
ABQ00274/standard; DNA; 24
XX
AC ABQ00274;
XX
DT 11-JUN-2002 (first entry)
XX
Oligonuclectide adapter/ca
XX
Oligonuclectide array; ada
XX
Oligonuclectide acidisequ
Array comprising adapter s
Array comprising adapter s 25-AUG-2000; 2000US-227948P 29-AUG-2000; 2000US-228854P Oligonucleotide array; adapter sequence; probe; Oligonucleotide adapter/capture probe 2001WO-US26519 (first entry) 24 ВP SS.

C C ၀ ၀

44444444

112 112 112 112 112 112 113 11.4

82585

112224444

ABQ00274 ABQ04512 ABQ04553 ABQ10840 ABQ10881 ABQ12376 ABQ12376 ABQ12417 AAQ35925 AAT95155

Oligonucleotide a Oligonucleotide a

ad add add add add

Human Human/monkey

or

monkey heavy

Array comprising adapter sequences useful for a target nucleic acid sequence, has different different specific capture probes

immobilizing or detecting
addresses comprising

Score

Length

В

IJ

Description

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Ş

1 CAAGGTCCAGNCC 13

Query Match Best Local Matches

l Similarity 12; Conserv

Conservative

85.7%; 92.3%; 5 C;

Score 12; DB 24; L Pred. No. 2.1e+03; Prematches 1;

Length 24; Indels

0,

Gaps

0

Sequence

24

BP;

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Α,

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ç;

6

Τ;

0

other;

nucleic acid.

Claim 1;

Page

50;

261pp;

English.

The invention relates to an oligonucleotide array (I) comprising at least 25 different addresses (adapter sequences) with each comprising a different capture probe selected from a group consisting of the sequences given in ABQ00010-ABQ13409. (I) is useful for immobilising a target nucleic acid sequence by attaching a adapter nucleic acid (ABQ00010-ABQ13409) to a target nucleic acid to form a modified target nucleic acid and contacting the modified target nucleic acid with (I). The steps of above method is useful for detecting a target nucleic acid, which further comprises detecting the presence of the modified target nucleic acid.

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RESULT 2
ABQ04512/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                B
                                 different addresses (adapter sequences) with each comprising a different addresses (adapter sequences) with each comprising a different capture probe selected from a group consisting of the sequences given in ABQ00010-ABQ13409. (I) is useful for immobilising a target nucleic acid sequence by attaching a adapter nucleic acid (ABQ00010-ABQ13409) to a target nucleic acid to form a modified target nucleic acid and contacting the modified target nucleic acid with (I). The steps of above method is useful for detecting a target nucleic acid, which further comprises detecting the presence of the modified target.
                                                                                                                                                                                                                                       Array comprising adapter sequences useful for a target nucleic acid sequence, has different
                                                                                                                                                                                                                                                                                                                                                                             25-AUG-2000; 2000US-227948P.
29-AUG-2000; 2000US-228854P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oligonucleotide adapter/capture probe 4503
                                                                                                                                                            The invention relates to an oligonucleotide array (I) comprising at least
                                                                                                                                                                                              Claim
                                                                                                                                                                                                                         different specific capture
                                                                                                                                                                                                                                                                                                                                               (ILLU-) ILLUMINA INC
                                                                                                                                                                                                                                                                                                                                                                                                                        27-AUG-2001; 2001WO-US26519
                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first
                                                                                                                                                                                              145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            array; adapter sequence; probe;
                                                                                                                                                                                           261pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
                                                                                                                                                                                                                         probes
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addresses comprising
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Sequence

24 BP;

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other

DX AC X

ABQ10840;

11-JUN-2002

(first entry)

RESULT 4 ABQ10840/c

ABQ10840 standard;

DNA;

24

BP.

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RESULT 3
ABQ04553
 В
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PR
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                                              Matches
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Best Local 9
                                                                                                                                                 25 different addresses (adapter sequences) with each comprising a different capture probe selected from a group consisting of the sequences given in ABQ00010-ABQ13409. (I) is useful for immobilising a target nucleic acid sequence by attaching a adapter nucleic acid (ABQ00010-ABQ13409) to a target nucleic acid to form a modified target nucleic acid and contacting the modified target nucleic acid with (I).
                                                                                                                                                                                                                                                                      different specific capture probes
                                                                                                                                                                                                                                                                                  Array comprising adapter sequences useful for a target nucleic acid sequence, has different
                                                                                             Sequence 24 BP;
                                                                                                                    nucleic
                                                                                                                                          The steps of above method is useful
                                                                                                                                                                                                                        The invention relates to an oligonucleotide array (I) comprising at least
                                                                                                                                                                                                                                                Claim 1; Page 145;
                                                                                                                                                                                                                                                                                                                    WPI; 2002-292068/33
                                                                                                                                                                                                                                                                                                                                           Gunderson
                                                                                                                                                                                                                                                                                                                                                                                         25-AUG-2000;
29-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200216649-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABQ04553;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABQ04553 standard;
                                                                                                                                which
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                            further comprises detecting the presence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19 CAAGGTCCAGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      l Similarity
12; Conserv
 σ
                    1 CAAGGTCCAGNCC
                                              12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 CAAGGTCCAGNCC 13
                                                                                                                    acid.
                                                          Similarity
CAAGGTCCAGACC
                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                         2000US-227948P
2000US-228854P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
                                                                                             σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      array; adapter sequence; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             adapter/capture
                                                                                             A.
                                                                                                                                                                                                                                             261pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry!
                                                          85.7%;
92.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    92.3%;
                                                                                             8 C; 5
                       13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24
                                                                                                                                                                                                                                                English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВP
                                                                                             G; 5 T; 0 other;
                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 12;
Pred. No.
                                                                    Score 12;
                                                          Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            probe
                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                          No.
                                                                                                                              for detecting a target nucleic acid, he presence of the modified target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4544
                                                         2.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.1e+03;
                                                                     DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BB
                                                                                                                                                                                                                                                                                  immobilizing or detecting
addresses comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 24
                                                                    Length 24
                                              Indels
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                                             0;
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                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                             0;
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RESULT 5
ABQ10881
ID ABQ1
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Best Local S
                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                           25 different addresses (adapter sequences) with each comprising at least different capture probe selected from a group consisting of the sequences given in ABQ00010-ABQ13409. (I) is useful for immobilising a target nucleic acid sequence by attaching a adapter nucleic acid (ABQ00010-ABQ13409) to a target nucleic acid to form a modified target nucleic acid and contacting the modified target nucleic acid with (I). The steps of above method is useful for detecting a target nucleic acid, which further comprises detecting the presence of the modified target nucleic acid.
                                                                                                            Synthetic
                                                                                                                                  Oligonucleotide
                                                                                                                                                      Oligonucleotide adapter/capture probe 10872.
                                                                                                                                                                                                    ABQ10881;
                                                                                                                                                                                                                          ABQ10881 standard;
                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 223; 261pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Array comprising adapter sequences useful for immobilizing or detecting a target nucleic acid sequence, has different addresses comprising different specific capture probes -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gunderson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-AUG-2000;
29-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-AUG-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligonucleotide array;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oligonucleotide
                                                                                       WO200216649-A2
                                                                                                                                                                              11-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ILLU-) ILLUMINA
         25-AUG-2000;
29-AUG-2000;
                                          27-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2002-292068/33
                                                                                                                                                                                                                                                                                  19
                                                                                                                                                                                                                                                                                                        1 CAAGGTCCAGNCC 13
                                                                                                                                                                                                                                                                                                                               12;
                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                  CAAGGTCCAGACC 7
                                                                                                                                                                                                                                                                                                                                                                        24 BP; 5 A; 5 C; 8 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                              Conservative
         2000US-227948P.
2000US-228854P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2000US-227948P
2000US-228854P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2001WO-US26519
                                          2001WO-US26519
                                                                                                                                                                             (first entry)
                                                                                                                                 array;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          adapter/capture probe 10831.
                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                     85.7%;
92.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      adapter sequence; probe;
                                                                                                                                 adapter
                                                                                                                                                                                                                           24
                                                                                                                                                                                                                           ВP
                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                      Score 12;
Pred. No.
                                                                                                                                 sequence; probe;
                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                   DB 24;
                                                                                                                                                                                                                                                                                                                                         .1e+03
                                                                                                                                 SS.
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                                                                                                                                                                                                                                                                                                                             0;
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RESULT 6
ABQ12376/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25 different addresses (adapter sequences) with each comprising a different capture probe selected from a group consisting of the sequences given in ABQ00010-ABQ13409. (I) is useful for immobilising a target nucleic acid sequence by attaching a adapter nucleic acid (ABQ00010-ABQ13409) to a target nucleic acid to form a modified target nucleic acid and contacting the modified target nucleic acid with (I). The steps of above method is useful for detecting a target nucleic acid, which further comprises detecting the presence of the modified target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Array comprising adapter sequences useful for immobilizing or detecting a target nucleic acid sequence, has different addresses comprising different specific capture probes
The invention relates to an oligonucleotide array (I) comprising at 25 different addresses (adapter sequences) with each comprising a different capture probe selected from a group consisting of the sequences in ABQ00010-ABQ13409. (I) is useful for immobilising a target
                                                                                                                 Array comprising adapter sequences useful for immobilizing or detecting a target nucleic acid sequence, has different addresses comprising
                                                                                                                                                                                        Gunderson
                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                            Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotide adapter/capture
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim
                                                                        Claim
                                                                                                   different
                                                                                                                                                           WPI; 2002-292068/33
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABQ12376;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABQ12376 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to an oligonucleotide array (I) comprising at least
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ILLU-) ILLUMINA INC
                                                                                                                                                                                                                     (ILLU-) ILLUMINA INC
                                                                                                                                                                                                                                                  29-AUG-2000;
                                                                                                                                                                                                                                                               25-AUG-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  further ic acid.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24 BP; 6 A;
                                                                                                   specific capture
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                2000US-227948P
2000US-228854P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                           2001WO-US26519
                                                                       242;
                                                                                                                                                                                                                                                                                                                                                                                                            array; adapter sequence; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         261pp;
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92.3%;
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C;
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                                                                       English.
                                                                                                  probes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                       probe 12367.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .1e+03;
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(!) is useful for

sequences

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RESULT 7
ABQ12417
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Db
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          δ
                                             Matches
                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                          The invention relates to an oligonucleotide array (I) comprising at least 25 different addresses (adapter sequences) with each comprising a different capture probe selected from a group consisting of the sequences given in ABQ00010-ABQ13409. (I) is useful for immobilising a target nucleic acid sequence by attaching a adapter nucleic acid to form a modified target nucleic acid and contacting the modified target nucleic acid with (I). The steps of above method is useful for detecting a target nucleic acid, with further comprises detecting the presence of the modified target nucleic acid.
                                                                                              Sequence
                                                                                                                                                                                                                                                     Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ABQ00010-ABQ13409) to a target nucleic acid to form a modified target nucleic acid to form a modified target nucleic acid to form a modified target nucleic acid with (1). The steps of above method is useful for detecting a target nucleic acid, which further comprises detecting the presence of the modified target
                                                                                                                                                                                                                                                                                   Array comprising adapter sequences useful for immobilizing or detecting a target nucleic acid sequence, has different addresses comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200216649-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oligonucleotide array; adapter sequence; probe; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oligonucleotide adapter/capture probe 12408
                                                                                                                                                                                                                                                                                                                                                                                               25-AUG-2000; 2000US-227948P
29-AUG-2000; 2000US-228854P
                                                                                                                                                                                                                                                                                                                                                                                                                                  27-AUG-2001; 2001WO-US26519
                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                       (ILLU-) ILLUMINA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABQ12417 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nucleic acid
                                                                                                                                                                                                                                                                                                                         2002-292068/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20
                       1 CAAGGTCCAGNCC 13
                                               12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 CAAGGTCCAGNCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12;
                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
CAAGGTCCAGACC 19
                                                                                              25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CAAGGTCCAGACC 8
                                                                                                                                                                                                                                                                            specific capture
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 5
                                               Conservative
                                                                                              BP; 6 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Α;
                                                                                                                                                                                                                                                   261pp;
                                                          85.7%;
92.3%;
                                                                                             8 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             85.7%;
92.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ū
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               c;
                                                                                             5 G;
                                                                                                                                                                                                                                                    English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8
                                                                                                                                                                                                                                                                          probes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ВP
                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 12;
Pred. No.
                                                           Pred. No.
                                                                    Score
                                                                                             6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7
                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               other;
                                                                     DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DΒ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .1e+03;
                                                          .1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24;
                                             ۲.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 25;
                                                                   Length 25;
                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                            0;
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ΧX
                          XXXX ID
                                                                                                                                   В
                                                                                                                                                           δÃ
                                                                                                                                                                                                                                            RESULT 9
AAT95155/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ΧX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 8
AAQ35925/c
ID AAQ35925 standard;
                                                                                                                                                                                          Matches
                                                                                                                                                                                                   Query Match
Best Local
                                                                 AAT95155 standard;
                                                                                                                                                                                                                                                                             families. The amplified sequences were cloned into p-Bluescript for sequencing. The amplified sequences could be used in the production of a recombinant antibody, comprising a human, chimpanzee or old world monkey immunoglobulin (Ig) constant region and an antigen (Ag) binding portion of an old world monkey Ig varible region, where the old world monkeys may be the same or different. The recombinant antibody may be used as a therapeutic agent for the treatment of
                                                                                                                                                                                                                                                                                                                                                                         The sequences given in AAQ35925-30 represent sense primers which used to amplify cynomolgus heavy chain variable region sequences. The amplified sequences belonged to a number of different gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Recombinant antibodies including Old World monkey portion and human portion - used for treatment of auto-immune diseases, infectious diseases, AIDS, tumours, diabetes, proliferative diseases, intestinal inflammations and allergies, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR; primer; region; lambda; human; constant; antigen; recombinant; antibody; chimpanzee; Ag; old world monkey; eczema; immunoglobulin; therapeutic; rheumatoid arthritis; p-Bluescript; ss.
                                                                                                                                                                                                                                            Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-JUL-1991;
23-MAR-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
             18-FEB-1998
                                                                                                                                                                                                                                                                   rheumatoid arthritis, eczema
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hanna
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24-JUL-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-FEB-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 W09302108-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human/monkey heavy chain variable region primer VH1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-JUN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ35925;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (IDEC-) IDEC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Amplify;
                                                                                                                                     17
                                                                                                                                                   1 CAAGGTCCAGNCCA 14
                                                                                                                                                                                       12;
                                                                                                                                                                                                   Similarity
                                                                                                                                   CCAGGTCCAGTCCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Newman RA,
                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                          BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                              Fig 8A;
            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PHARM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      91US-0735064.
92US-0856281.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92WO-US06194
                                                                 DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                            92pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chain reaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
                                                                                                                                                                                                                                          5 C;
                                                                                                                                                                                                   81.4%;
85.7%;
                                                                 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17
                                                                                                                                                                                                                                          9
                                                                                                                                                                                                                                                                                                                                                                                                                              English.
                                                                 ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВP
                                                                                                                                                                                                                                          ე:
                                                                                                                                                                                       0,
                                                                                                                                                                                                   Pred. No.
                                                                                                                                                                                                          Score 11.4;
                                                                                                                                                                                                                                                                   and immunological disorders
                                                                                                                                                                                                                                          3 T;
                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                          0 other;
                                                                                                                                                                                                   .6e+03;
                                                                                                                                                                                                                 DB 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            light; chain; variable;
                                                                                                                                                                                                                Length
                                                                                                                                                                                       Indels
                                                                                                                                                                                                                 17;
                                                                                                                                                                                       0;
                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                     were
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Human or monkey Ig heavy chain variable region primer VH1

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10-JUL-1992;
25-JUL-1991;
23-MAR-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                rheumatoid arthritis; lupus erythematosus; PCR primer; systemic lupus erythematosus; SLE; Hashimoto's thyroiditis; multiple sclerosis; myasthenia gravis; type 1 diabetes; uveitis; nephrotic syndrome; psoriasis; atopical dermatitis; contact dermatitis, eczematous dermatitides; seborrheic dermatitis; Lichen planus; Pemplugus; bullous pemphigus; Epidermolysis bullosa; urticaria; angioedemas; vasculitides; erythema; cutaneous eosinophilia; alopecia areata;
                                                                                                                                                                                                                                                             constant region coding sequence.

The chimeric Ab may be used as a passive or active therapeutic for B cell lymphoma, infectious, e.g. AIDS, autoimmune and inflammatory diseases and transplant rejection. The Ab can be used in its native form or as part of an Ab/chelate, Ab/drug or Ab/toxin complex. Additionally, a whole Ab, or a fragment (Fab2, Fab, Fv) may be used as an imaging reagent or as a potential vaccine or immunogen in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence is a PCR primer for a human or monkey immunoglobulin heavy chain variable region cDNA, which was used in the development of a novel method for producing a chimeric antibody (Ab), comprising the variable region of an Old World monkey (OWM) Ab and the constant region of a human Ab. The method comprises amplifying DNA from the OWM using a primer complementary to the 5' leader sequence of the OWM Ab gene and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Production of humanised antibodies containing monkey variable region - used for the treatment of B cell lymphoma, autoimmune disease, inflammatory disease and infection, e.g. by HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
Cebus apella.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           reversible obstructive airways disease; intestinal inflamma; intestinal allergy; Coeliac disease; proctitis; eosinophiligastroenteritis; mastocytosis; Crohn's disease; ulcerative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             inflammatory disease; transplant rejection; imaging reagent; vaccine; immunogen; immunotherapy; anti-idiotypic response; immunosuppression; treatment; prevention; proliferative disease; immunologically mediated disease; hyperproliferative disease; immunologically mediated disease;
                                                                                                                                  active immunotherapy to generate anti-idiotypic responses An anti-CD4 recombinant Ab, or a fragment can also be used induce immunosuppression and can therefore be used to tree prevent resistance to, or rejection of transplanted organs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-JAN-1995;
07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US5681722-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         human; variable region; chimeric antibody;
B cell lymphoma; infectious disease; AIDS; autoimmune disease;
inflammatory disease; transplant rejection; imaging reagent;
                                                                                                         tissues,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1997-535052/49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Old World monkey; immunoglobulin; Ig; heavy chain; CD4;
                                hyperproliferative diseases, cutaneous manifestations of immunologically mediated diseases (e.g. rheumatoid arthr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    fusing the resulting variable region coding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          related allergy; migraine; rhinitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IDEC
                                                                                                         autoimmune,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Newman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PHARM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      92US-0912292.
91US-0735064.
92US-0856281.
95US-0379072.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 95US-0478039
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                95US-0478039
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RΑ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    84pp; English
                                                                                              inflammatory, proliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Raab
erythematosus, Hashimoto's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  eczema;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the OWM Ab gene and sequence with a human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nal inflammation;
eosinophilia;
                                                                                                     and
                                                                                                                                                               be used to
                            arthritis,
                                                                                                                                      organs
                                                                                                                                         or
S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  colitis
                                lupus
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Qy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 10
AAT91566/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                        25-JUL-1991;
23-MAR-1992;
25-JAN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 dermatitis, contact dermatitis and further eczematous dermatitides, seborrheic dermatitis, Lichen planus, Pemplugus, bullous pemphigus, Epidermolysis bullosa, urticaria, angioedemas, vasculitides, erythema, cutaneous eosinophilias, alopecia areata), reversible obstructive airways disease, intestinal inflammations and allergies (e.g. Coeliac disease, proctitis, eosinophilia gastroenteritis, mastocytosis, Crohn's disease and ulcerative
                              A new chimeric antibody (Ab) has been developed comprising a human or chimpanzee immunoglobulin (Ig) constant region and an Old World monkey antigen (Ag)-binding region. An anti-CD4 antibody (Ab) comprising the light and heavy-chain variable regions encoded by AAT91564 and AAT9156
                                                                                           Example 1; Fig
                                                                                                                                                                     WPI;
                                                                                                                                                                                                                   (IDEC-)
                                                                                                                                                                                                                                                                                                                                                                                                                            Anti-CD4 antibody; cynomolgus monkey; immunoglobulin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cynomolgus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          half-life and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         colitis) and food related allergies (e.g. migraine,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         thyroiditis, multiple sclerosis, myasthenia gravis, type 1 diabetes, uveitis, nephrotic syndrome, psoriasis, atopical dermatitis, contact dermatitis and further eczematous derm
                                                                                                                   antigen-binding region
                                                                                                                                                                                            Hanna
                                                                                                                                                                                                                                                                                                       25-JUL-1991;
                                                                                                                                                                                                                                                                                                                              19-AUG-1997
                                                                                                                                                                                                                                                                                                                                                      US5658570-A
                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                      antigen
                                                                                                                                                                                                                                                                                                                                                                                                                   chimpanzee;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-JAN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT91566 standard; DNA; 17
                                                                                                                                                                                                                                                                             10-JUL-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       induce a human anti-Ab response. It also has a relatively
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ab of the invention is non-immunogenic and therefore does
                                                                                                                                                                     1997-424183/39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 CAAGGTCCAGNCCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       l Similarity
12; Conserv
                                                                                                                                                                                                                    IDEC
                                                                                                                                                                                                                                                                                                                                                                                                     binding
                                                                                                                               antibodies for human therapy - coee immunoglobulin constant region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CCAGGTCCAGTCCA 4
                                                                                                                                                                                           Newman RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                     monkey
                                                                                                                                                                                                                                                                                                                                                                                                                   chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 3
                                                                                                                                                                                                                   PHARM CORP
                                                                                                                                                                                                                                        92US-0912292.
91US-0735064.
92US-0856281.
95US-0379072.
                                                                                            ,,
                                                                                                                                                                                                                                                                                                       91US-0735064
                                                                                                                                                                                                                                                                                                                                                                                                     neric antibody; human
region; tumour; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          full effector
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A.
                                                                                            46pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                    heavy chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   81.4%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 11.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No. 4.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   variable region primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          function with human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                     therapy; 0. polymerase
                                                                                                                               comprising human or on and an Old World
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18;
                                                                                                                                                                                                                                                                                                                                                                                                     ; Old Worl
ase chain
                                                                                                                                                                                                                                                                                                                                                                                                               World
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cells and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           rhinitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                       VH1.
                                                                                                                                                                                                                                                                                                                                                                                                     reaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    long
                                                                                                                               monkey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and
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                                                                                                                                                                                                                                                                                                                                                                                                     SS
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is also new. The present sequence represents a primer, VH1, used in the amplification of cynomolgus monkey immunoglobulin heavy chain regions. The Abs are useful for human therapy, especially of tumours. Old work

used in the

AAT91565

Old world

human therapy,

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RESULT 11
AAT62902/
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                                                                                     Query Match
Best Local
                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
                                                                                                                                                          Sequences are provided for heavy chain variable regions VH1-VH6 (AAT62902-07), heavy chain constant region IgM and IgG1-4 antisense strands (AAT62908-09), kappa and lambda light chain variable regions (AAT62910-11) and kappa and lambda light chain constant region antisense strands (AAT62912-13). Novel monoclonal and chimaeric antibodies (see also AAM14924) comprise heavy and light chain variable regions (AAM1492-33) from macaque anti-human CD4 monoclonal antibody E9.1 and human gamma-1 or gamma-4 (AAW14925-27) constant
                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         monkeys are sufficiently different from humans to allow Abs against human Ags, even relatively conserved Ags such as CD4 and CD54, to be raised in these monkeys, and are sufficiently similar to humans to avoid host anti-Ab immune responses when the Abs are introduced into humans.
                                                                                                                                                                                                                                                                             Disclosure; Fig 8; 155pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        VH1 heavy chain
                                                                                                                                                                                                                                                                                                                      useful
                                                                                                                                                                                                                                                                                                                                  constant
                                                                                                                                                                                                                                                                                                                                             Chimeric antibody
                                                                                                                                                                                                                                                                                                                                                                      WPI; 1997-201913/18.
                                                                                                                                                                                                                                                                                                                                                                                                                       (IDEC-) IDEC PHARM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                06-SEP-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9709351-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          transplant rejection; HIV; therapy; CE9.1; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cynomolgus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CD4; monoclonal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAT62902;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT62902 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2D4; monoclonal antibody; chimaeric antibody; recombinant antibody; cynomolgus monkey; autoimmune disease; rheumatoid arthritis; leukaemia; lymphoma; graft-versus-host disease; asthma;
į,
                     17 CCAGGTCCAGTCCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17
                                              1 CAAGGTCCAGNCCA 14
                                                                         12;
                                                                                                                                                                                                                                                                                                                   for treatment of autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    l Similarity
12; Conser
                                                                                   Similarity
                                                                                                                           17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CAAGGTCCAGNCCA 14
                                                                                                                                                                                                                                                                                                                                  domains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CCAGGTCCAGTCCA 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17 BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                              Newman RA,
                                                                                                                          ₿₽;
                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                95US-0523894
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        96WO-US14324
                                                                                                                          w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       variable region.
                                                                                                                                                                                                                                                                                                                 comprising monkey variable domains and human - affects CD4-mediated immune functions, espent of autoimmune disease, e.g. rheumatoid
                                                                                                                          ?
                                                                                  81.4%;
85.7%;
                                                                                                                          5 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               81.4%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 C; 6
                                                                                                                                                                                                                                                                                                                                                                                               Reff ME;
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                                                                                                                          σ
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                                                                                                                          G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>ن</u>
                                                                      Score 11.4; In Pred. No. 4.6e 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pred. No. 4.6
); Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3 T; 0 other;
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                                                                                                                         0 other;
                                                                                   6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .6e+03;
                                                                                             DB 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 17;
                                                                       Indels
                                                                                              Length
                                                                                                                                                                          monoclonal
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AAV23793/c
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AAV31430/c
                                                                                                                                                                                                Best Loc
Matches
                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                      25-JUL-1991;
23-MAR-1992;
10-JUL-1992;
25-JAN-1995;
                                                                                                                                                                                                                                                         This primer is used for the PCR amplification of the heavy chain variable (VH) region of cynomolgus immunoglobulin sequence. This is used in the construction of a chimeric human-monkey anti-CD4 antibody. The antibody comprises antigen-binding sequences from an Old World monkey antibody and human constant domain sequences. The antibody specifically binds to CD4 and can be used in the treatment of rheumatoid or psoriatic arthritis.
        rheumatoid arthritis; PCR primer;
                 Anti-CD4 antibody; Old World monkey;
                                                                                     AAV23793;
                                                                                                                                                                                                                                      Sequence 17
                                                                                                                                                                                                                                                                                                                                              administration of chimeric human-monkey anti-CD4 antibody
                                                                                                                                                                                                                                                                                                                                                         Treatment of rheumatoid and psoriatic arthritis
                                                                                                                                                                                                                                                                                                                                                                              WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US5756096-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Anti-CD4 antibody;
monkey; rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cynomolgus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAV31430;
                                               Primer
                                                                  29-JUL-1998
                                                                                                         AAV23793
                                                                                                                                                                                                                                                                                                                            Example 1; Fig 8; 47pp; English
                                                                                                                                                                                                                                                                                                                                                                                                  Hanna
                                                                                                                                                                                                                                                                                                                                                                                                                   (IDEC-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Macaca cynomolgus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-AUG-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAV31430 standard; DNA;
                                                                                                                                                                                                         Local
                                                                                                                                                                                                                                                                                                                                                                              1998-321461/28
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                                                                                                                                                          17
                                                                                                                                                                                                l Similarity
12; Conser
                                                                                                                                                                            1 CAAGGTCCAGNCCA 14
                                                                                                                                                                                                                                                                                                                                                                                                                   IDEC
                                                                                                        standard; cDNA;
                                                                                                                                                         CCAGGTCCAGTCCA 4
                                                                                                                                                                                                                                                                                                                                                                                                Newman RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immunoglobulin
                                                                                                                                                                                                                                       BP;
                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                   PHARM CORP
                                                                  (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                     95US-0476237.
91US-0735064.
92US-0856281.
92US-0912292.
95US-0379072.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 95US-0476237
                                                                                                                                                                                                                                       ω
                constant domain; eczema;
                                                                                                                                                                                                                                       A; 5 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antigen-binding; treatment; chimeric; human;
arthritis; psoriatic arthritis; PCR primer;
                                                                 entry)
                                             antibody coding sequence
                                                                                                                                                                                                       81.4%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                 Raab
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17
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                                                                                                                                                                                               Score 11.4; I
Pred. No. 4.66
0; Mismatches
                                                                                                                                                                                                0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      region amplifying
                                                                                                                                                                                                                                       w
                                                                                                                                                                                                                                       Η,
         SS
                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                       other;
                                                                                                                                                                                                         4.6e+03;
                                                                                                                                                                                                                   DВ
                          variable heavy
                  immuno-modulated disease;
                                                                                                                                                                                                                  19;
                                             fragment
                                                                                                                                                                                                2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primer
                                                                                                                                                                                                                 Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       VH1
                                                                                                                                                                                                                    17;
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AAV05683/c
ID AAV05683
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This sequence is a PCR primer for DNA encoding an anti-CD4 antibody (Ab). The amplified sequence can be used in the method of the invention for treating a subject, where the treatment comprises administration of an Ab. The, method comprises the administration of an antibody which has an Old World monkey (e.g. baboon or macaque) variable region which binds to an antigen (Ag) (or Ag binding portion), and a human constant domain. The method is useful for the treatment of eczema and immuno-modulated diseases and especially rheumatoid arthritis. The recombinant antibodies used are sufficiently different from native monkey antibodies to allow human antigens to raise these antibodies, but similar enough to human antibody so there is no immune response to the antibodies in humans. Compared to antibodies used in therapy in prior art, these antibodies do not induce human anti-antibodies on repeated administration. They also have longer half-lives and do not have a lack of effector function with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic. Primate sp:
                                                                                                                                                                                                                                  macaque; immunoglobulin; heavy chain variable region; chimpanzee; hybrid;
lymphoma; AIDS; autoimmune disease; inflammatory disease; transplant;
                                                                                                                                                                                                                                                                                                         Primer; PCR; amplification; leader sequence; human; monkey; baboon
                                                                                                                                                                                                                                                                                                                                                                                        Human/monkey
                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Improved method for antibody treatment - uses an antibody comprising an Old World monkey variable region and a human constant domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1998-296690/26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hanna
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (IDEC-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-MAR-1992;
05-DEC-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-JUL-1992;
25-JUL-1991;
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                                                                                                                       Synthetic.
                                           Macaca sp.
                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-MAY-1998
   Papio sp
                                                                                                                                                                                              tumour; antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       'n
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12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CCAGGTCCAGTCCA 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fig
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                    heavy chain variable region primer VH1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PHARM CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      92US-0856281.
95US-0379072.
95US-0476349.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8
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91US-0735064
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Α,
                                                                                                                                                                                                  SS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               81.4%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Raab
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 11.4; DB 19;
Pred. No. 4.6e+03;
D; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4; DD
. 4.6e+03;
. 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 15
AAT43427/c
ID AAT43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.0-JUL-1992;
25-JUL-1991;
23-MAR-1992;
25-JAN-1995;
07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antigen-binding region; and (ii) a human or chimpanzee Ig constant region. The hybrid antibodies can be used as passive or active therapeutic agents against human diseases, e.g. B-cell lymphoma, AIDS, autoimmune and inflammatory diseases, transplant rejection or tumours, or for producing therapeutic and diagnostic conjugates. Although evolutionary distant monkeys are used to raise antibodies against human antigens, they are sufficiently similar that they produce antibodies similar to human antibodies, such when the monkey antibodies are introduced into humans, no host anti-antibody response is stimulated.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US5693780-A
   15-MAY-1995;
                                                                           14-MAY-1996;
                                                                                                                                                      21-NOV-1996
                                                                                                                                                                                                                                                                                                                                                                            animal model; pig; monoclonal antibody; prime
polymerase chain reaction; phage display; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of a recombinant antibody comprising: (i) an Old World monkey Ig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This sequence represents a sense primer derived by comparing sequences from human and monkey (especially baboon (Papio sp.) and macaque (Macaca sp.)) immunoglobulin (Ig) heavy chain regions. The primer is used to amplify the monkey heavy chain variable region for construction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleic acid encoding hybrid antibody - comprising antigen-binding region and human or chimp constant % \left( 1\right) =\left( 1\right) +\left( 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1998-031820/03.
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                                                                                                                                                                                                                                  WO9636358-A1
                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                        Xenograft rejection; xenotransplantation; organ
animal model; pig; monoclonal antibody; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       VH1 family specific primer for human heavy chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-FEB-1997 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT43427 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-DEC-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CCAGGTCCAGTCCA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 3 A; 5 C; 6 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 92US-0856281.
95US-0379072.
95US-0481869.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8; 46pp; English.
95US-0440621
                                                                           96WO-US06804
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91US-0735064
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      81.4%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 11.4; DB 19 Pred. No. 4.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 19;
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Indels Length

0;

Gaps

0,

transplant;
PCR;

밁 Q monkey

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Search completed: June 21, 2003, 22:26:29 Job!time: 78.7812 secs
                                                                         В
                                                                                                       δÃ
                                                                                                                                       Query Match 81.4
Best Local Similarity 85.7
Matches 12; Conservative
                                                                                                                                                                                                                                           PCR primers VH1-VH6 (AAT43427-32) are Ig family specific heavy chain 5' primers designed for the amplification of antibody VH region cDNA and for the introduction of cohesive termini compatible with directional ligation into the unique Noti or Spel sites of the SurfZAP vector. They were used with 3' primers in the PCR amplification of antibody heavy chain variable region cDNA from lymphocytes of a human transplant patient who had received a porcine xenograft. A human IgM heavy chain, human IgM light chain Fab library was generated using phage display and panned for anti-donor xenograft antibody sequences.
                                                                                                                                                                                                               Sequence 18 BP; 4 A; 5 C; 6 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Inhibiting xenograft rejection by modifying antigen expression of the graft - prevents binding of anti-donor antibody and prolongs graft survival
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cramer DV, Makowka L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1997-011852/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 16; Page 121; 135pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CEDA-) CEDARS SINAI MEDICAL CENT.
                                                                   1 CAAGGTCCAGNCCA 14
| |||||||| |||
17 CCAGGTCCAGTCCA 4
                                                                                                                                                          81.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wu G;
                                                                                                                                             0,
                                                                                                                                         Score 11.4; DB 18;
Pred. No. 4.6e+03;
0; Mismatches 2;
                                                                                                                                                                           Length 18;
                                                                                                                                           Indels
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                                                                                                                                         Gaps
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Minimum DB
Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Database :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Title:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OM nucleic - nucleic search, using sw model
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       Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score
       seq length: 0 seq length: 50
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1: /cgn2_6/ptodata/1,

2: /dgn2_6/ptodata/1,

3: /cgn2_6/ptodata/1,

4: /dgn2_6/ptodata/1,

5: /dgn2_6/ptodata/1,

6: /cgn2_6/ptodata/1,
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1 CAAGGTCCAGNCCA 14
       441362 seqs, 153338381 residues
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Copyright (c) 1993 - 2003 Compugen Ltd
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/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/FQTUS_COMB.seq:*
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       DB
US-08-379-072A-5
US-08-478-039-51
US-08-478-039-51
US-08-476-249A-51
US-08-476-237-1
US-08-523-894-44
US-09-523-487-13
US-09-523-487-13
US-08-646-687-3
US-08-646-687-3
US-08-646-687-3
US-08-478-039-20
US-08-478-039-20
US-08-478-039-20
US-08-646-687-3
US-08-814-111-59
US-09-287-145A-59
US-09-287-145A-52
US-09-184-658-24
US-09-184-658-24
US-09-184-658-24
US-09-184-658-24
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258.254 Million cell updates/sec
  sequence
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Sequence
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	10	10	10	10	10	10	10.4.	10.4	10.4	10.4	10.4	10.4	10.4	10.4	10.4	10.4	10.4	10.4
	71.4	71.4	71.4	71.4	71.4	71.4	74.3	74.3	74.3	74.3	74.3	74.3	74.3	74.3	74.3	74.3	74.3	74.3
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ALIGNMENTS	US-08-379-078-314	US-08-379-081B-314	US-09-746-694-40	3 US-09-289-267-65	US-07-841-652-8	L US-07-767-135-17	3 US-08-589-939-61	5 5427925-17	5 5187077-19	2 US-08-924-440-4	FCT-US94-10957-19	us-09-225-928-653	US-08-859-998-653	US-09-462-136-10	US-08-691-814B-53	US-09-487-792-50	US-09-819-964-16	US-08-949-155-16
	Sequence 314, App	Sequence 314, App	Sequence 40,.Appl	Sequence 65, Appl	Sequence 8, Appli	Sequence 17, Appl	Sequence 61, Appl	Patent No. 5427925	Patent No. 5187077	Sequence 4, Appli	Sequence 19, Appl	Sequence 653, App	Sequence 653, App	Sequence 10, Appl	Sequence 53, Appl	Sequence 50, Appl	Sequence 16, Appl	Sequence 16, Appl

## ALIGNMENTS

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STRANDEDNESS: single;
TOPOLOGY: linear;
MULECULE TYPE: DNA (ger
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                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/0
FILING DATE: 25-VAN-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: P.O. BOX 1404
CITY: Alexandria
STATE: Virginia
COUNTRY: United States
ZIP: 22313-1404
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                            INFORMATION FOR SEQ ID NO:
                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: (703) 836-6620 TELEFAX: (703) 836-2021
                                                                                                                                                                                                                                    APPLICATION NUMBER: US 07/735,064
FILING DATE: 25-JUL-1991
ATTORNEY/AGENT INFORMATION:
NAME: Rea, Teresa Stanek
                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0:
FILING DATE: 23-MAR-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: RECOMBINANT ANTIBODIES FOR HUMAN THERAPY NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                NAME: Rea, Teresa Stanek
REGISTRATION NUMBER: 30,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0,
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RAAB, Ronald W.
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10-JUL-1992
NN DATA
                 .DNA (genomic)
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Patent No.
                              Matches
                                           Query Match
Best Local :
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Best Local Similarity
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                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 35,030
REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
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MEDIUM TYPE: Floppy disk
                                                                                                                 POSITION IN GENOME:
                                                                                                                                                TOPOLOGY: 1
                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                  ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                     FILING DATE: 25-JUL-1991
ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: US 07/912,292
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                                                                                                                                                           STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 699 Prince CITY: Alexandria
                                                                                                    CHROMOSOME/SEGMENT:
                                                                                                                                                                                                                                                                  TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 07/735,064 FILING DATE: 25-JUL-1991
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1 CAAGGTCCAGNCCA 14
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                                            Similarity
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                                                                                                                                                                                          nucleic acid
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                                                                                                                                                                                                                                                      703-836-2021
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                             Conservative
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                                           81.4%;
85.7%;
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85.7%;
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                                                                                                    heavy chain variable region
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                          Score 11.4; D
Pred. No. 5.8e
O; Mismatches
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                                           8e+02
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                                                          DB 1;
                                                       Length 17;
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                           Indels
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RESULT 4
US-08-476-349A-51/c
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                                                            Sequence 51, Application US/08476349A Patent No. 5750105
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5 Sequence 5, Appitus
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GENERAL INFORMATION:
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Best Local S
                             GENERAL INFORMATION:
APPLICANT: Newman
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                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0.
FILING DATE: 23-MAR-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: 'PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                 TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 536
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: (703) 836-6620
                 APPLICANT:
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 APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE: 10-JU
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ZIP: 223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/0 FILING DATE: 25-JAN-1995
                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER:
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                                                                                                                                                         17 CCAGGTCCAGTCCA
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RY: United States
22313-1404
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                                                                                                                                                                                                                                                                                                                                                                                                           (703) 836-2021
Hanna, Nabil
Raab, Ronald W.
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RAAB, Ronald W
                             Newman,
                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                  linear
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                              Roland A.
                                                                                                                                                                                                                                   81.4%;
85.7%;
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                                                                                                                                                                                                                                  Score 11.4; DB 1, Pred. No. 5.8e+02
                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                    Length 17;
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                                                                                                                                                                                   Sequence
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                                                                                                                                                                   Patent No.
                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                           ANTI-SENSE: NO
POSITION IN GENOME:
CHROMOSOME/SEGMENT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 25-JAN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/912,292
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Rel CURRENT APPLICATION DATA:
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ADDRESSEE: BURNS, DOANE,
                                                                                                     APPLICANT: NEWMAN, Roland
APPLICANT: HANNA, Nabil
APPLICANT: RAAB, Ronald W.
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TELECOMMUNICATION INFORMATION:
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ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER:
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APPLICATION NUMBER: US 08/379,072
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: . 10-JUL-1992
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                           STREET:
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                                                                                                                                                                                                                                                                                           1 CAAGGTCCAGNCCA 14
            SSEE: Burns, Doane, Swecker & Mathis
T: 'P.O. Box 1404
Alexandria
                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                 Application US/08476237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Teskin Esq., Robin L.
Virginia
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                                                                                                                                                                                                                                                                                                                                     81.4%;
85.7%;
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Pred. No. 5
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Best Local :
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INFORMATION FOR SEQ ID NO:
                               APPLICATION NUMBER: US/08
FILING DATE: 06-SEP-1995
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Teskin, Robin L.
                                                                                                                                                                        ZIP: 22314-3187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 25-JUL-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                              SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
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CURRENT APPLICATION DATA:
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                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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NAME: Teskin, Robin L.
REGISTRATION NUMBER: 35,030
REFERENCE/DOCKET NUMBER: 012712-165
                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                           CITY:
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                                                                                                                                                                                                                                                                       699 Prince Street
                                                                                                                                                                                                                            USA
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Pred. No. 5.8e+02;
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SOFTWARE: FRSTSEQ for Windows Version 3.0
SEQ ID NO 13
LENGTH: 22
                                                                                                                                                                                    Sequence 13, Appli
Patent No. 6232098
                                                                                                                                                                                                                                                                                                                                                  Matches
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Best Local :
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                                  CURRENT APPLICATION NUMBER: US/09/523,487
CURRENT FILING DATE: 2000-03-10
                                                                                                               APPLICANT: Conklin, Darrell APPLICANT: Foster, Donald C. APPLICANT: Gao, Zeren
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/320,095
                                                                    FILE REFERENCE:
                                                                                TITLE OF INVENTION: KUNITZ DOMAIN POLYPEPTIDE AND MATERIALS TITLE OF INVENTION: AND METHODS FOR MAKING IT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ¤ARLIER APPLICATION NUMBER: US 60/087,032
¤ARLIER FILING DATE: 1998-05-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: KUNITZ DOMAIN POLYPEPTIDE AND MATERIALS TITLE OF INVENTION: AND METHODS FOR MAKING IT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Conklin, Darrell C.
APPLICANT: Foster, Donald C.
APPLICANT: Gao, Zeren
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
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 FILING DATE:
                    APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                    1 CAAGGTCCAGNCCA 14
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1999-05-26
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85.7%;
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BER: 09/320,095
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SOFTWARE: Fai
SEQ ID NO 13
FRNGTH: 22
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Best Local :
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APPLICANT: Newman
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                                                                                                                          TELEFAX: 703-836-2021 INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 08/379,072
PRILING DATE: 25-JAN-1995
                                                                                                                                                                      NAME: Teskin Esq., Robin L.
REGISTRATION NUMBER: 35,030
REFERENCE/DOCKET NUMBER: 01.
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 22313-1404
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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ANTI-SENSE:
ORIGINAL SO
                               MOLECULE TYPE:
                                                                                                           SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                     FILING DATE: 25-JUL-1991
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 0 FILING DATE: 10-JUL-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0,
                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/0 FILING DATE: 07-JUN-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Alexandria
                                               TOPOLOGY:
                                                                                                                                                         TELEPHONE:
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                                                                                            CENGTH:
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12; Conserv
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                                                                          nucleic acid
SOURCE
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                                                                                         26 base pairs
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Raab, Ronald W
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                                              linea
                                                                                                                                                           703-836-6620
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                                                                                                                                                                                                                                                                                                     23-MAR-1992
                                                             single
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85.7%;
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                                                                                                                                                                                        012712-160
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Pred. No. 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Version #1.30
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CHROMOSOME/SEGMENT: US-08-476-349A-20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                    NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: PC-L
SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 2222
COMPUTER READABLE FORM:
MPDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Homo sapiens or Monkey POSITION IN GENOME:
                               POSITION IN GENOME
                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: 703-836-6620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: RECOMbinant Antibodies for Human Therapy
NUMBER OF SEQUENCES: 114
CORRESPONDENCE ADDRESS:
ADDRESSEE: BURNS, DOANE, SWECKER & MATHIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                            ORIGINAL SOURCE:
                                                                                                                                                                     SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                              FILING DATE: 25-JUL-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Newman,
                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
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                                                                           ANTI-SENSE:
                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 0:
FILING DATE: 25-JAN-1995
                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US 0' FILING DATE: 10-JUL-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 07 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
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                                            ORGANISM:
                                                                                                                       STRANDEDNESS:
                                                                                                                                                                                                       TELEFAX:
                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
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                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                REGISTRATION NUMBER:
                                                                                                                                                      ENGTH:
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                                                                                                                                       nucleic acid
                                                                                                                                                                                                                                                                               Teskin Esq., Robin L
                                                                                                                                                   26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             699 Prince St.
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Raab, Ronald W
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                                                                                                        linear
                                            Homo sapiens or Monkey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IBM PC compatible SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                              NUMBER: US 07/856,281
23-MAR-1992
                                                                                           CDNA
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             heavy chain primers with SalI site
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RESULT 12
US-08-646-367-3/c
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Patent No. 6136310
                                                                                                                             Query Match
Best Local Similarity
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Best Local (
                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 22314-3187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
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INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                       STRANDEDNESS:
TOPOLOGY: lin-
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-836-6620
TELEFAX: 703-836-2021
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ATTORNEY/AGENT INFORMATION:
NAME: Teskin, Robin L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
                                                                                                                                                                                                  POSITION IN GENOME
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Teskin, Robin L.
REGISTRATION NUMBER: 35
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE: 06-SE
                                                                                                                                                                                  CHROMOSOME/SEGMENT: VH1 leader sequence
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                                                                                                                                                                                                                 ORGANISM:
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                                                        26 CCAGGTCCAGTCCA 13
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                                                                                    1 CAAGGTCCAGNCCA 14
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                                                                                                                Conservative
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                                                                                                                                                                                                                Human or Monkey
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                                                                                                                                                                                                                                                         DNA (genomic)
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                                                                                                                              81.4%;
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85.7%;
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                                                                                                                             Score 11.4;
Pred. No. 6
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                                                                                                              Mismatches
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                                                                                                                             6e+02;
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                                                                                                              Indels
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                                                                                                              Gaps
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Sequence 3, Application US/08646367 Patent No. 5959085 GENERAL INFORMATION: APPLICANT: Pierre Garrone

APPLICANT:

Pierre Garrone Odile Djossou

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RESULT 13
US-08-373-190-59/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: MARASCO, WAYNE
APPLICANT: HASELTINE, WILLIAM
TITLE OF INVENTION: METHOD OF INTRACELLULAR BINDING OF PROTEINS
NUMBER OF SEQUENCES: 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 908-298-5388
                                            COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOTTWARE: FASTSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/373,190
FILLING DATE: 17-JAN-1995
                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 27 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: 908-298-2987
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STREET: Kenilworth
CITY: Kenilworth
STATE: New Jersey
                 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Microsoft Word 5.1a
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Macintosh 7.5.3
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                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
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 APPLICATION NUMBER:
                                   CLASSIFICATION:
                                                                                                                                                                                        COUNTRY: US
ZIP: 02109
                                                                                                                                                                                                                           STATE:
                                                                                                                                                                                                                                         STREET: 130
CITY: BOSTON
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Foulke, Cynthia L.
REGISTRATION NUMBER: 32,364
REFERENCE/DOCKET NUMBER: SF0403K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: FILING DATE: May 16
                                                                                                                                                                                                                                                                           ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24 CCAGGTCCAGTCCA 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 CAAGGTCCAGNCCA 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  l Similarity
12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleic acid
                                                                                                                                                                                                                           ΜA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E: Schering-Plough Corporation 2000 Galloping Hill Road
                                                                                                                                                                                                                                                         130 WATER STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Francois Fossiez
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                           DIKE, BRONSTEIN, ROBERTS & CUSHMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Diskette
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                81.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Against Human Cytokines And Methods Of Making And Using Such Antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human Monoclonal Antibodies
PCT/US93/06735
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 11.4;
Pred. No. 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 27;
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US-08-438-190A-59
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US-08-438-190A-59/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 59, Application US/08438190A Patent No. 5965371
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 617-523-6440
TELEX: STRE UR 2002
INFORMATION FOR SEQ ID NO:
                                                                              TELEFAX: (617) 523-644
TELEX: 200291 STRE UR
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HYPOTHETICAL: NO ANTI-SENSE: NO FRAGMENT TYPE: ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: DAVID G. CONLIN; DIKE, BRONSTEIN, ROBERTS ADDRESSEE: CUSHMAN
                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 523-3400
                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: EISENSTEIN, RONALD I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION: TELEPHONE: 617-523-3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: METHOD OF INTRACELLULAR BINDING TITLE OF INVENTION: PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: MARASCO, WAYNE APPLICANT: HASELTINE, WILL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS: LENGTH: 30 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 16-JUL-1993 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: LE BOSTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                        NAME: EISENSTEIN, REGISTRATION NUMBER:
                              STRANDEDNESS:
                                                TYPE:
                                                                                                                                                    TELEPHONE:
                                                                                                                                                                                        REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/438,190A
                                                                                                                                                                                                                                                                                                                                                                 COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: US
                                                                   LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27 CCAGGTCCAGTCCA 14
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                          H: 30 base pairs nucleic acid DEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  MASSACHUSETTS
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                                                                                                                                    (617) 523-6440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HASELTINE, WILLIAM
                                                                                                                                                                                                                                                                                                                                              IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                 linear
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Pred. No. 6.1e+02
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Search completed: June 22, 2003, 00:01:38 Job time: 18.625 secs
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US-08-803-085-14/c
                                                                                                                                                                                           ; MOLECULE TYPE: DNA (genomic) US-08-803-085-14
                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/803,085
FILING DATE: 20-FEB-1997
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Teskin, Robin L.
REGISTRATION NUMBER: 35,030
REFERENCE/DOCKET NUMBER: 012712-353
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 836-2021
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 14, Patent No. 6
                                                                                                                             Matches
                                                                                                                                            Query Match
Best Local Similarity
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: REFF, Mitchell E.
APPLICANT: KLOETZER, William S.
APPLICANT: NAKAMURA, Takehiko
TITLE OF INVENTION: GAMMA-1 ANTI-HUMAN CD23 MONOCLONAL
TITLE OF INVENTION: ANTIBODIES AND USE THEREOF AS THERAPEUTICS
RUMBER OF SEQUENCES: 35
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: Virginia
COUNTRY: United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: P.O. Box 1404
CITY: Alexandria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
                                                            1 CAAGGTCCAGNCCA 14.
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                                                                                                                             Conservative
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                                                                                                                                           81.4%;
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Pred. No. 6.1e+02;
0; Mismatches 2;
                                                                                                                                            Score 11.4;
Pred. No. 6
                                                                                                                             Mismatches
                                                                                                                                           6.1e+02;
                                                                                                                                                           DB 3;
                                                                                                                                                     Length 30;
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Maximum DB
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Maximum Match 100%
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                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No.
                                                                                                                                                                                                                                                                                                                  Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       seq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      | length: 0
| length: 50
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'Match Length
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11111987654321
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Gapop 10.0 , Gapext 1.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /cgn2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/US08_BUBCOMB.seq:*
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/cgn2_6/ptodata/2/pubpna/US60_PUBCOMB.seq:*
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US-10-067-790-16

US-10-067-892-16

US-10-067-893-16

US-10-067-893-16

US-10-067-893-16

US-10-024-018-2

US-10-227-1218-38

US-10-215-112-10872

US-10-215-157-13
                                                                                                                                                                                                    0 US-09-964-666-14

0 US-09-964-412-14

US-09-940-185-281

US-09-940-185-4261

US-10-211-357-44
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                 US-10-103-686-14
US-09-019-441-14
                                                  US-09-850-165-42
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Sequence 14, Appl
Sequence 14, Appl
Sequence 281, App
Sequence 4261, App
Sequence 44, Appl
Sequence 11, Appl
Sequence 16, Appl
Sequence 16, Appl
Sequence 16, Appl
Sequence 2, Appli
Sequence 2, Appli
Sequence 13, Appl
Sequence 13, Appl
Sequence 13, Appl
Sequence 13, Appl
Sequence 14, Appl
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																	74.3					81.4	81.4	81.4	81.4	81.4
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ALIGNMENTS .	)98-263B-1	-098-263в-8	2-86(	US-10-098-263B-62978	098-263в-6	098 - 2	0-098-263B-	N 3	US-10-098-263B-35907	US-10-098-263B-35398	- 1	US-09-864-636A-2607	308	US-10-215-112-6573	US-09-906-032A-3	US-10-087-325-19	US-09-864-636A-2609	US-09-801-274-1163	US-10-098-263B-45863	US-09-971-980-21	US-09-930-423-2985	-825-805-	-805-	-805-	US-09-925-664-52	US-09-825-805-219
	102		15		6251		Sequence 57464, A				Sequence 35397, A	2607	Sequence 10, Appl	657	Sequence 3, Appli			Sequence 1163, Ap	4586	1, A	Sequence 2985, Ap		987,	954,	52,	Sequence 219, App

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US-09-964-667-14
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                                                                                                                                                          INFORMATION FOR SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/09/964,667
FILING DATE: 28-eep-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein &
STREET: 1100 New York Ave., Suite 600
TOPOLOGY: linear
MOLECULE TYPE: cDNA
SEQUENCE DESCRIPTION: SEQ ID NO: 14:
                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 14 base pairs
                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 0609.4370000
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
TELEFAX: 202-371-2540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Transgenic Animals and Cell Lines Screening Drugs Effective for the
                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
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Wands, Jack R.
                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                        NAME: Esmond, Robert W. REGISTRATION NUMBER: 32,893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Washington STATE: DC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of Alzheimer's Disease
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e Treatment '
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of Alzheimer's Disease

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RESULT 3
US-09-964-412-14
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                                                                       Sequence 14, Application US/09964412 Patent No. US20020129391A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 14, Application US/09964666 Patent No. US20020104108A1 GENERAL INFORMATION:
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Best Local
                                                                                                                                                                                                                                                                                        Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 14:
TITLE OF INVENTION: Transgenic Animals and Cell Lines for Screening Drugs Effective for the Treatment or Prevention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
TELEFAX: 202-371-2540
                                        APPLICANT: de la Monte, Suzanne Wands, Jack R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox, P.L.L.C.
STREET: 1100 New York Ave., Suite 600
                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                          SEQUENCE DESCRIPTION: SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
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Wands, Jack R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                   1 CAAGGTCCAGNCCA 14
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                                                                                                                                                                                                 CAAGGTCCAGNCCA 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Esmond, Robert W.
REGISTRATION NUMBER: 32,893
REFERENCE/DOCKET NUMBER: 06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/09/964,666 FILING DATE: 28-Sep-2001 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
ZIP: 20005-3934
                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
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Pred. No. 5.3e+02;
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5.3e+02;
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Query Match
Best Local Similarity
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                                                                                         ; OTHER INFORMATION: Computer Generated Probe Sequence US-09-940-185-281
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US-09-940-185-281/c
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                                                                                                                                                               NUMBER OF SEQ ID NOS: 4768
SOFTWARE: Patentin version 3.1
SEQ ID NO 281
LENGTH: 24
TYPE: DNA
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Publication No. US20030096239A1
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Probes and Decoder Oligonucleotides FILE REFERENCE: A 69605-1 CURRENT APPLICATION NUMBER: US/09/940,185 CURRENT FILING DATE: 2001-08-27
                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 60/227,948
PRIOR FILING DATE: 2000-08-25
PRIOR APPLICATION NUMBER: US 60/228,854
PRIOR FILING DATE: 2000-08-29
                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Gunderson, Kevin
                                                                                                                                           ORGANISM: Artificial Sequence
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COMPUTER READABLE FORM:
MEDLUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: PAtentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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MOLECULE TYPE: cDNA
SEQUENCE DESCRIPTION: SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler,
STREET: 1100 New York Ave.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: <Unknown> ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/09/964,412 FILING DATE: 28-Sep-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                  Chee, Mark
                   Conservative
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                                  85.7%;
92.3%;
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Pred. No.
                                  Score 12; DB 9;
Pred. No. 1.8e+03;
                   Mismatches
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. 5.3e+02;
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                 Indels
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CAAGGTCCAGNCC

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RESULT 6
US-10-211-357-44/c
Sequence 44, Application US/10211357
Publication No. US20030077275A1
GENERAL INFORMATION:
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Best Local Similarity
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LENGTH: 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Publication No. US20 GENERAL INFORMATION:
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CURRENT FILING DATE: 2001-08-27
PRIOR APPLICATION NUMBER: US 60/227,948
PRIOR FILING DATE: 2000-08-25
PRIOR APPLICATION NUMBER: US 60/228,854
PRIOR FILING DATE: 2000-08-29
PRIOR FILING DATE: 2000-08-29
NUMBER OF SEQ ID NOS: 4768
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Gunderson, Kevin
APPLICANT: Chee, Mark
TITLE OF INVENTION: Probes and Decoder Oligonucleotides
FILE REFERENCE: A-69605-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn version 3.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESSEE: BURNS, DOANE, SWECKER & MATHIS
STREET: 699 Prince Street
                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                          PRIOR APPLICATION DATA:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Hanna, Nabil
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               APPLICATION NUMBER: US/09/612,914A
FILING DATE: 10-Jul-2000
APPLICATION NUMBER: US 08/523,894
FILING DATE: 06-SEP-1995
                                                                                                                        APPLICATION NUMBER: US/10/211,357
FILING DATE: 05-Aug-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                        STATE: VA
                                                                                                                                                                                                                                                                                                                                                                                             CITY: Alexandria
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92.3%;
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Mitchell E.
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Pred. No. 1.8e+03;
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US-09-850-165-1/c
                                                                     Query Match
Best Local S
Matches 12
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LENGTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/850,165
CURRENT FILING DATE: 2001-05-08
PRIOR APPLICATION NUMBER: 09/082,472
PRIOR FILING DATE: 1998-05-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: NEWMAN, ROLAND A.
                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 114
                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 07/PRIOR FILING DATE: 1991-07-25
                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 07/856,281
                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 1995-04-17
PRIOR APPLICATION NUMBER: 07/912,292
PRIOR FILING DATE: 1992-07-10
                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 08/476,237 PRIOR FILING DATE: 1995-06-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: RECOMBINANT ANTIBODIES FOR HUMAN THERAPY
                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 1992-03-23
                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 08/397,072
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: 037003-0280614
                                                                                                                                                                             FEATURE:
                                                                                                                                                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                                                                 TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 44:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CHROMOSOME/SEGMENT: VH1 heavy chain variable region SEQUENCE DESCRIPTION: SEQ ID NO: 44:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ANTI-SENSE: NO
POSITION IN GENOME:
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LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: DNA (genomic)
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   17
                                                                     l Similarity
12; Conserv
                                  1 CAAGGTCCAGNCCA 14
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CCAGGTCCAGTCCA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: 703-836-6620
TELEFAX: 703-836-2021
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RAAB, RONALD W.
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85.7%;
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85.7%;
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                                                                                     Score 11.4;
Pred. No. 46
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Pred. No. 4
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RESULT 8 US-10-067-790-16/c ; Sequence 16, Application US/10067790

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; LOCATION: ()..()
; OTHER INFORMATION: primer
US-10-067-892-16
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; LOCATION: ()..()
; OTHER INFORMATION: primer
US-10-067-790-16
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Best Local Similarity
Matches 12; Conserv
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Best Local
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                                                                      Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Publication No. US20030035807A1
                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/1
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APPLICANT: TURPEN, THOMAS ANTIGEN VACCINES FOR TREATING B CELL LYMPHOMAS AND OTHER CAN
FILE REFERENCE: 18696-169194
                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US/09/522,900 PRIOR FILING DATE: 2000-03-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS:
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                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: McCORMICK, Alison
APPLICANT: TUSE, Daniel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ENERAL INFORMATION:
                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ublication No. US20030039659Al
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            l0-067-892-16/c
                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: SELF ANTIGEN VACCINES FOR TREATING B CELL LYMPHOMAS TITLE OF INVENTION: AND OTHER CANCERS THE REFERENCE: 18696-169194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Unknown
                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                     LENGTH: 18
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15 CCAGGTCCAGTCCA 2:
                               1 CAAGGTCCAGNCCA 14
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                                                                                     Similarity
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REINL, Stephen
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TURPEN, Thomas
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85.7%;
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                                                                                   Score 11.4;
Pred. No. 4
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; OCATION: ()..()
; OTHER INFORMATION: primer
US-09-539-382-16
                                                                                                 ; OTHER INFORMATION: primer US-10-067-893-16
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LENGTH: 18
TYPE: DNA
ORGANISM: Unknown
FEATURE:
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                                                                                                                                                                                 SOFTWARE: PatentIn version 3.0
SEQ ID NO 16
LENGTH: 18
TYPE: DNA
ORGANISM: Unknown
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                                 Matches
                                                 Query Match
Best Local Similarity
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APPLICANT:
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APPLICANT: TUSE, Daniel
APPLICANT: REINL, Stephen
                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 60/155,579 PRIOR FILING DATE: 1999-09-24 NUMBER OF SEQ ID NOS: 62
                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/067,893
CURRENT FILING DATE: 2002-02-08
PRIOR APPLICATION NUMBER: 09/522,900
PRIOR FILING DATE: 2000-03-10
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CURRENT FILING DATE: 2000-03-10
PRIOR APPLICATION NUMBER: US 60/155,579
PRIOR FILING DATE: 1999-09-24
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APPLICANT: TURPEN, Thomas
TITLE OF INVENTION: SELF ANTIGEN VACCINES FOR TREATING B CELL LYMPHOMAS AND
FILE REFERENCE: 18696-169195
                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: TURPEN, Thomas
TITLE OF INVENTION: SELF ANTIGEN VACCINES FOR TREATING B CELL LYMPHOMAS AND OTHER
FILE REFERENCE: 18696-169194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: McCORMICK, Alison
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                                                                                                                                                                      FEATURE:
                                                                                                                                 NAME/KEY: misc_feature
LOCATION: (). ()
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1 CAAGGTCCAGNCCA 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TUSE, Daniel
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                                                 81.48;
85.78;
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                                                 Pred. No.
                                                               Score 11.4;
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Pred. No. 4
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RESULT 14
US-10-215-112-10872
Sequence 10872, Application US/10215112
Publication No. US20030082596A1
GENERAL INFORMATION:
APPLICANT: Michael Mittmann
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                                                                                                                                                                                                                                                                                                                  ; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Primer US-09-927-121B-38
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; ORGANISM: Homo sapiens
US-10-024-018-2
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US-10-024-018-2
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Best Local S
Matches 12
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LENGTH: 22
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APPLICANT: GOLD, DANIEL P.
APPLICANT: SHOPES, ROBERI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2, Application US/10024018 Publication No. US20030004140A1
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Publication No.
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Best Local :
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TITLE OF INVENTION: METHOD AND COMPOSITION FOR ALTERING A B CELL MEDIATED TITLE OF INVENTION: PATHOLOGY PATHOLOGY FILE REFERENCE: 032077,0003

CURRENT APPLICATION NUMBER: US/09/927,121B

CURRENT FILING DATE: 2001-08-10
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TITLE OF INVENTION: METHODS FOR MODULATING CELL-ADHESION MEDIATED DRUG
FILE REFERENCE: 215101.01403
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT FILING DATE: 2001-08-10 NUMBER OF SEQ ID NOS: 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 09/795,474
PRIOR FILING DATE: 2001-03-01
PRIOR APPLICATION NUMBER: US 60/186,199
PRIOR FILING DATE: 2000-03-01
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CURRENT FILING DATE: 2001-12-21
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TITLE OF INVENTION: Method of Genetic Analysis of Probes:
                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Artificial Sequence
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12; Conserv
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85.7%;
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Pred. No. 3
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Pred. No. 3.
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3.9e+03;
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                                                                                                                                                                                                                                                                             Length 22
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RESULT 15
US-10-211-357-13/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 10872
LENGTH: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 13, Application US/10211357 Publication No. US20030077275A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/215,112
CURRENT FILING DATE: 2002-08-08
NUMBER OF SEQ ID NOS: 14936
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Hann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                             TELEPHONE: 703-836-66
TELEFAX: 703-836-2021
INFORMATION FOR SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
                                                                                                                                                                                                                                                                                                                      FILING DATE: 10-Jul-2000
APPLICATION NUMBER: US 08/523,894
FILING DATE: 06-SEP-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: 703-836-6620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: BURNS, DOANE,
                                     MOLECULE TYPE:
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
                    ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Recombinant Anti-CD4 Antibodies for Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10 CTAGGTCCAGACCA 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12;
                                                                                                TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                               NAME: Teskin, Robin L. REGISTRATION NUMBER: 35,030
                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 05-Aug-2002
CLASSIFICATION: <Unknown>
                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/09/612,914A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/10/211,357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 22314-3187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: VA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Alexandria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 699 Prince Street
ORGANISM: Human or Monkey
                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 012712-165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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Reff, Mitchell E.
                                                              DNA
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85.7%;
                                                          (genomic)
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Pred. No. 3.8e+03;
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POSITION IN GENOME:

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CHROMOSOME/SECHENT: VH1 leader sequence Us-10-211-357-13

Query Match
Best Local Similarity 81.4%; Score 11.4; DB 9; Length 26;
Best Local Similarity 85.7%; Pred. No. 3.8e-03;
Matches 12; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 CAMGGTCCAGNCCA 14

Db 26 CAMGGTCCAGNCCA 13

Search completed: June 22, 2003, 03:18:25

Job:time: 43.3438 secs
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Title:
Perfect score:
Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                   Post-processing: Minimum Match 0%
| Maximum Match 100%
| Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                         Database :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Minimum DB seq length: 0
Maximum DB seq length: 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Run on:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OM nucleic - nucleic search, using sw model
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                                                                                                                                                    14:
16:
17:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      June 21, 2003, 21:28:37; Search time 604.734 Seconds (without alignments) 374.936 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16154066 seqs, 8097743376 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IDENTITY_NUC Gapop 10.0 , Gapext 1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GenCore version 5.1.6 Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CAAGGTCCAGNCCA 14
                                                                                                                                                                                            gb_est1: *
gb_est2: *
gb_htc: *
gb_est3: *
gb_est4: *
gb_est5: *
em_estom:*
gb_gss:*
em_gss_inv:*
em_gss_inv:*
em_gss_inv:*
em_gss_fvn:*
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em_gss_fwn:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

		d				
Result		Query				
No.	Score	Score Match	Score Match Length DB ID	BB	ij	Description
c 1	11.4	11.4 81.4		10	BE282036	BE282036 601102010
2	11.4	81.4	50	10	AV428742	AV428742 AV428742
ω	11	78.6	50	14	H55190	H55190 CHR220129 C
c 4	10.4	74.3	19	17	AZ825396	AZ825396 2M0100N04
c 5	10.4	74.3	31	17	AZ437960	AZ437960 1M0226A14
6	10.4	74.3	46	13	BI908094 ·	BI908094 603067183

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	ω	7
	9.4					9.4			9.4									٠			,													10		10	10	10.4
67.1	67.1	67.1	67.1	67.1	7.	7.	7.	7.	67.1	0	0	0	0	0	0	0	0	0	0	0	0.	0	0	0	0.	0	0.	μ.	71.4	۳	1.	71.4	71.4	71.4	71.4	71.4	71.4	74.3
37	37	34	33	32	30	23	23	23	22	50	50	50	50	49	49	49	48	47	43	40	39	38	32	29	28	28	25	50	50	50	50	46	44	42	35	34	20	50
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81520	523	103472	Z7913	79316	45629	78545	48067	30782	8950		AU106942						248579		9054		5916	30F0	9029	9256	53A0	AZ327470	1395					æ	m	19	(4)	66941	81	AU107443
 15205 wk72c11.	180523 uc70a08	J034722 BJ0347	791392 2M004j	Z793160 2M0046	456295 1M0259	Z785457 2M0029	Z480676 1M0302	307822 1M0010	89506 1M015	794 yy16g04	AU10694	06721 AU10672	03167 AŪ10316	57977 tj54d07	87238 oj35q08	6193 nx25d08	85794 1M0313K2	9 602937	790549 SALK_057	880280 vv99h07.r		L464113 T. brúce	Z990297 2M0273N2	92566 1M0326F1	72207 T. bruce	0 1M0050J1	13950 1M050	5659 AU10565	3657	2547 AU10254	02541 AU1025	86886 Human exon	89056 1M0319G	48919 yb07c07.s1	F036425	6941 ua38c02.r	Z819886 2M0091	AU107443 AU107443

## ALIGNMENTS

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Email: cgapbs-r@mail. Tissue Procurement: G CDNA Library Prepara CDNA Library Arrayed DNA Sequencing by: I Clone distribution: found through the I.M http://image.llnl.gov Plate: LLAMB643 row. Plate: LLAMB643 row. High quality sequence	JOURNAL Unpublished (1999) COMMENT Contact: Robert St	S E	×	ACCESSION BE282036 VERSION BE282036.1 GI:9157330 KEYWORDS EST. SOURCE house mouse.	RESULT 1 BE282036/C BC282036 LOCUS BE282036 BE28206 B
Email: cgapbs-r@mail.nih.gov Tissue Procurement: Gilbert Smith, Ph.D. cDNA Library Preparation: Life Technologies, Inc. cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: LLAM8543 row: c column: 04 High quality sequence start: 2 High quality sequence stop: 49.	Unpublished (1999) Contact: Robert Strausberg, Ph.D.	1 (bases 1 to 49) NIH-MGC http://mgc.nci.nih.gov/. NATional Thetitute of Health Mammalian Game Collection (MGC)	Mus musculus Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.	:I:9157330	BE282036 49 bp mRNA linear EST 26-OCT 2000 601102010F1 NCI_CGAP_Lu29 Mus musculus cDNA clone IMAGE:3494331 5′,

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RESULT
H55190
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AV428742
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Matches 12
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Best Local S
Matches 12
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MEDLINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AUTHORS
TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM
                                                                                                                                                                                                         COUNT
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              ω
                                                              17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lotus japonicus.

Lotus japonicus

Lotus japonicus

Lotus japonicus

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;

Spermatophyta; Magnoliophyta; Fabaceae; Papilionoideae; Loteae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AV428742

50 bp mRNA linear AV428742 Lotus japonicus young plants (two-week old) japonicus cDNA clone MWM035d10_r 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                       Kazusa DNA Research Institute
Yana 1532-3, Kisarazu, Chiba 292-0812, Japan
Email: asamizu@kazusa.or.jp, URL:http://www.kazusa.or.jp/en/plant/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 CAAGGTCCAGNCCA 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12;
                                                                                          1 CAAGGTCCAGNCCA 14
                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Erika Asamizu
The First Laboratory for Plant Gene Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                            legume, Lotus japonicus
DNA Res. 7 (2), 127-130 (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Asamizu, E., Nakamura, Y., Sato, S. Generation of 7137 non-redundant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AV428742.1 GI:7678124
EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       japonicus cDNA
AV428742
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                                                                                                                                          Similarity
                                                           CAAGGTCGAGTCCA 30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (bases 1 to 50)
                                                                                                                           Conservative
                                                                                                                                                                                                         22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                 /organism="Lotus japonicus"
/db_xref="taxon:34305"
/db_xref="taxon:34305"
/clone="MWH035d10_r"
/clone_lib="Lotus japonicus young plants (two-week old)"
/dev_stage="young plants (two-week old)"
/dev_stage="young plants (two-week old)"
/note="Vector: pbluescriptil SK-; Site_1: EcoRI; Site_2:
xhoI; isolate=Miyakojima MG-20"

xhoI; isolate=Miyakojima MG-20"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note="organ: lung; Vector: pCMV-SPORT6; Site_1: SalI; Site_2: NotI; Cloned unidirectionally. Primer: Oligo of Library constructed by Life Technologies. Investigator providing samples: Gilbert Smith, NIH"

a 11 c 27 g 9 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /tissue_type="spontaneous tumor, metastatic to mammary.
Stem cell origin."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone="IMAGE:3494331"
/clone_lib="NCI_CGAP_Lu29"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Mus musculus"
/strain="C2ECH II"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                    location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /lab_host="DH10B"
                                                                                                                                         81.4%;
85.7%;
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                                                                                                                           0;
                                                                                                                                       Score 11.4; DB 10
Pred. No. 4.2e+04;
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Pred. No. 4.2e+04;
                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and Tabata, S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           expressed
                                                                                                                                                       DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence
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Unpublished (2000)
Contact: Robert B. W
University of Utah G
University of Utah
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H55190.1
                                                                                     and Wright, D., Weiss, Mouse whole genome s
                                                                                                                                                       1 (bases 1 to 19)
Dunn, D., Aoyagi, A.,
                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                  2M0100N04R Mouse 10kb plasmid clone UUGC2M0100N04 R, DNA sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Building 149, 13th Tel: 6177249616
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Molecular Neurogenetics Unit
Massachusetts General Hospital
Building 149, 13th St., Charlestown MA 02129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 50)
Trofatter, J.A., Lo
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                       plasmid inserts
                                                                                                       Islam,H., Longacre,S., Mahmoud,M.,
,M., Rose,M., Rose,R., Stokes,R.,
and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                  Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and Buckler, A.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                  house mouse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: buckler@helix.mgh.harvard
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone_lib="Chromosome 22 exon"
/lab_host="E. coli DH5a"
/note="Vector: pBluescriptIIKS+;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /db_xref="taxon:9606"
/clone="C22_166"
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91.7%;
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                    Genome
                                      Weiss
                                                                                     scaffolding
                                                                                                                                                           Barber, M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22
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                                                                                                                                                                                                                                                                                                                      DNA sequence.
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d UUGC1M library
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                                                                                                                      Tingey, A.,
                                                                                                                                          Meenen, E.,
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; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                                      20-FEB-2001
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RESULT 5
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                                                                                                                                                                                                                                                                                                                                                                                                   1M0226A14R Mouse 10kb plasmid UUGCIM library clone UUGCIM0226A14 R, DNA sequence.
AZ437960
                                                                                                                                  Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
and Wright,D.,Welss,R.
                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 31)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
                                                                                                                                                                                                                                                                                                                                                         AZ437960.1 GI:10561973 GSS.
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Location/Qualifiers
University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                          Mus musculus
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Insert Length: 10000 Std Erro
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                                            Contact: Robert B.
                                                                  Unpublished (2000)
                                                                                                                   Mouse whole genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AZ437960
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                                                                                             inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWDA2 (gil4732114 gblnAF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coll XL10-Gold (Stratagene) cells and selected for ampicallin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="Mus musculus
/strain="C57BL/6J"
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                                                 Weiss
                                                                                                               scaffolding with paired
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                                                                                                                                                                                    Meenen, E., Pedersen, T., Reilly
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                                                                                                                 reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15 AAGCTCCAGCCCA 3
                                                                                           Contact: Robert Strausberg, Ph:D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies,
                                                                                                                                                                                  1 (bases 1 to 46)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Fax: 801 585 7177
                                                                                                                                                                 Unpublished (1999)
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cDNA Library Preparation: Life Technologies, Inc. CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732.114 [gb] AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coll XL10-Gold (Stratagene) cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and selected for ampicillin resistance 8 \text{ c} = 10 \text{ g} = 5 \text{ t}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0226A14"
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NBLAN134NF,
AU107443
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                                                                                                                                                                                                                           Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki.y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sug, S. Construction and characterization of a full length-enriched of 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 (bases 1 to 50)
Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J., Sese,J., Hai, H., Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo,K., Sakaki,Y., Nakamura,Y., Suyama,A. and Sugano,S.
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                                                                                                                                                                                                                                                                                                                                                                                                                           Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
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                                                   /clone_lib="Sugano Homo sapiens cDNA library"
/note="Differential display comparison of unt
dimethylfumarate treated U937 cells"
a 11 c 17 g 13 t
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/tissue_type="leukocyte"
/lab_host="DH10B"
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/db_xref="taxon:9606"
/clone="IMAGE:5216252"
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/db_xref="taxon:9606"
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and Wright,D.,Weiss,R.
Mouse whole genome scaffolding wi
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Fax: 801 585 7177
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University of Utah Genome Center
University of Utah
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Insert Length: 10000 Std Error: 0.00
Plate: 0091 row: H column: 16
Seg primer: CACACAGGAAACAGCTATGACC
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                                                                                                                  (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWAP2 (gil4732114 j@blAF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and sheared adaptored mouse DNA was annealed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
                                     adaptored vector DNA, and transformed into chemically-competent E. coli XLLO-Gold (Stratagene) and selected for ambicillin resistance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                  purified. The sheared, adaptored mouse DNA was annealed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M0091H16"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                           Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /lab_host="E. Coli strain XL10-Gold, T1-resistant,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Mus musculus"
selected for ampicillin resistance. 3 \text{ c} 4 \text{ g} 6 \text{ t}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 84.6%;
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0; Mismatches 2;
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71.48;

Score 10;

DB 17;

Length 20;

Вр QΥ

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REFERENCE
AUTHORS
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AA966941/c
LOCUS
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ORIGIN
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                                          Query Match
Best Local S
Matches 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Marra,M., Hillier,L., Allen,M., Bowles,M., Dietrich,N., Dubuque,T., Geisel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., Morris,M., Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B., Theising,B., Wylie,T., Lennon,G., Soares,B., Wilson,R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 AAGGTCCAGNC 12
       1 CAAGGTCCAG 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Trace considered overall poor quality Possible reversed clone: similarity on wise primer -28ml3 rev2 ET from Amersham High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WashU-HHMI Mouse EST Project
Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Marra M/Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The WashU-HHMI Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Waterston, R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CHAIN, MITOCHONDRIAL PRECURSOR ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ua38c02.rl Soares_mammary_gland_NbMMG Mus musculus cDNA clone
IMAGE:1348994 5' similar to SW:ATPD_RAT P35434 ATP SYNTHASE DELTA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AA966941.1 GI:3140834
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                                                                Similarity
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                                          Conservative
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a 11 c
                                                                                                                                                                         RNA provided by Dr. Minoru constructed and normalized
                                                                                                                                                                                                            T 3']; double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector.
                                                                                                                                                                                                                                                                  /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="IMAGE:1348994"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eutheria;
1 to 34)
                                                                                                                                                                                                                                                                                         RI; 1st strand cDNA was primed with a Not primer [5'
                                                                                                                                                                                                                                                                                                       /note="Organ: mammary gland; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: EcoRI; 1st_strand cDNA was primed with a Not I - oligo(dT)
                                                                                                                                                                                                                                                                                                                                                                                   /tissue_type="mammary gland"
/dev_stage="4 weeks"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                 /lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                          /sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                     'clone_lib="Soares_mammary_gland_NbMMG"
                                                            71.4%;
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                                            0;
                                                            Score 10;
Pred. No.
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0; Mismatches 1;
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                                                                                                                                                                           Bento Soares and M. Fatima
                                                                                                                                                                                            Wayne State Univ.
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                                                                             Length 34;
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CDNA clone
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                                          0;
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                                        Gaps
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                                          0;
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AUTHORS
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T48919/c
LOCUS
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ORGANISM
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TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           10;
 l (bases 1 to 42)
Hillier,L., Lennon,G., Becker,M., Bonaldo,M.F., Chiapelli,B.,
Chissoe,S., Dietrich,N., DuBuque,T., Favello,A., Gish,W., Haw
,M., Hultman,M., Kucaba,T., Lacy,M., Le,M., Le,N., Mardis,E.,
,B., Morris,M., Parsons,J., Prange,C., Rifkin,L., Rohlfing,T.
                                                                                                                                                                                                                                                                                                                                                                                                                    4 GGTCCAGNCCA 14
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BF036425
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                                                                                                                                                                           EST
                                                                                                                                                                                            T48919.1
                                                                                                                                                                                                                                           yb07c07.sl Stratagene placenta (#937225) Homo sapiens cDNA clone IMAGE:70476 3' similar to similar to gb:M17885 60S ACIDIC RIBOSOMAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
Plate: LLAM9604 row: d column: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tissue Procurement: DCTD/DTP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Robert Strausberg, Ph.D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mammalia; Eutheria;
1 (bases 1 to 35)
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                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                             PROTEIN PO (HUMAN),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                 GGTCCAGGCCA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CAAGGTCCAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /tissue_type="adenocarcinoma"
/lab_host="DH10B (phage-resistant)"
/note="Organ: ovary; Vector: pCMV-SPORT6; Site_1: Not1;
Site_2: Sal1; Cloned unidirectionally. Primer: Oligo d
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/clone_lib="NIH_MGC_66"
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/db_xref="taxon:9606"
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V., Mardis, E., Moore
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                                                              University of Utah Genome Center University of Utah
                                                                                                                          Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
                                                                                                                                                                  Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Du Islam,H., Longacre,S., Mahmoud,M., Meenen,E., M., Rose,M., Rose,R., Stokes,R., Tingey,A.,
                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                         AZ489056 44 bp DNA linear GSS 0: 1M0319G12R Mouse 10kb plasmid UUGCLM library Mus musculus clone UUGClM0319G12 R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This clone is available royalty-free through LLNL; IMAGE Consortium (info@image.llnl.gov) for further Trace considered overall poor quality
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Other_ESTs: yb07c07.rl
Contact: Wilson RK
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                                                                                             Contact: Robert B.
                                                                                                                                                          and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                  Mus musculus
                                                                                                                                                                                                                                                                                                               AZ489056.1 GI:10658433
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/clone="IMAGE:70476"
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/db_xref="GDB:491373"
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83.3%;
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1.2e+05;
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Meenen,E., Pedersen,T., Rei
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3 (bases 1 to 46) Ohira, M.
                                                                                                                                                                                                                                                                                                                                                                                                    GSS; exon trapping for Down's syndrome region Homo sapiens DNA, clone:E20-10.
                                     Unpublished
                                                                                                              Ohki, M.
                                                                                                                              Ohira,M.,
                                                                                                                                                                        Genome Res. (1997) In press
                                                                                                                                                                                                       chromosome 21
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                                                                                                                                                                                                                                                 and Ohki, M.
                                                                                                                                                                                                                                                                   Ohira, M., Seki, N., Nagase, T., Hattori, M., Sakaki, Y., Eki, T.,
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Mammalia; Eutheria;
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D86886
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynuclectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114 [gb] AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, "adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coll xL10 Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0319G12"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="Vector:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /lab_host="E. Coli strain XL10-Gold, T1-resistant,
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Y., Saito,T.,
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., Ichikawa,H.
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                                                                                                                                                                                                                                                                                                           Suzuki, r., Yoshitomo-Nakagawa, K., Maruyama, K., Suyama, A. and Sugano, S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Yutaka Suzuki
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/note="Differential display comparison of untreated and
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Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and
S. Construction and characterization of a full length-enrich
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Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J., Sese,J., Hata,H., Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo,K., Sakaki
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Department of Virology
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/note="Differential display comparison of untreated and dimethylfumarate treated U937 cells"
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# ALIGNMENTS

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Matobba,R., Okubo,K., Hori,N., Fukushima,A. and Matsubara,K. The addition of 5' coding information to a 3'-directed cDNA library improves analysis of gene expression	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  1 (bases 1 to 49)	Homo sapiens	Homo sapiens Male cell_line:HepG2 cDNA to mRNA, clone_lib:Kiseru.	gene signature.	D17183.1 GI:598688	D17183	Human HepG2 3' region MboI cDNA, clone hmd2h04m3.	HUMD2H04M3 49 bp mRNA linear PRI 04-FEB: 1999		

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RESULT 3
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de la Monte, S. and Wands, J.
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/db_xref="taxon:9606"
/sex="Male"
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/clone_lib="Kiseru"
15 c 12 g
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from patent US 5830670.
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Pred. No. 3.3e
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Pred. No. 2.1e+04;
D; Mismatches 7
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Sequence 63
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AR073178.1
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27 TTCAGCCTGGGTGACAGAGCGAGATCT 1
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                                              Schork,N. and Skierczynski,B.
Methods of genetic cluster analysis and
Patent: WO 0129257-A 48 26-APR-2001;
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                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Patent: US 5948888-A 63 07-SEP-1999;
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/organism="Homo sapiens"
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AX411689
AX411689.1 GI:21444223
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Sequence 7 from Patent WO0228897
AX411691
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                                                                                                                            Deleersnijder,W., Blockx,H. and de Moor,L
Human gyprotein coupled receptor and uses
Patent: WO 0228897-A 7 11-APR-2002;
SOLVAY PHARMACEUTICALS B V (NL)
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/note="Primer"
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/db_xref="taxon:32630"
/note="Primer"
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24
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AX466859
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AX056763
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Patent: WO 0075319-A 29 14-DEC-2000;
REGENERON PHARMACEUTICALS, INC. (US)
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1 (bases 1 to 38)
Papadopoulos,N.J., Davis,S. and Yano
Modified chimeric polypeptides with
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synthetic construct
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                                                                                                              /note="reverse primer"
5 c 6 g
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/db_xref="taxon:32630"
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/db_xref="taxon:32630"
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Cargill,M., Ireland,J.S. and Lander,E.S.
                                                                                                                      N
                                                                                                                                                                                                                                                                                                                                    artificial sequences.

1 (bases 1 to 38)

Papadopoulos,N.J., Davis,S. and Yancopoulos,G

Modified chimeric polypeptides with improved
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Patent: WO 0166800-A 8 13-SEP-2001,
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH (US)
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ON PHARMACEUTICALS, INC. (US)
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25 from Patent WO0075319
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/db_xref="taxon:9606"
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99138837
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Klonowski.K.D., Primiano, L.L. and Monestier, M.
Atypical VH-D-JH rearrangements in newborn autoimmune MRL mice 3. Immunol. 162 (3), 1566-1572 (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19;
   Direct Submission Submitted (12-JUN-1998) Microbiology and Immunology, Temple
                             2 (bases 1 to 50)
Monestier, M. and Klonowski, K.
                                                                                                                                                      Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unclassified
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Pastan, I.H., Kreitman, R.J. and Puri, R.
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Matches 19
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                                                    Gasche, C., Zakeri, S.M. and Reinisch, W. Mammalian interleukin-10 (il-10) receptor variants Patent: Wo 0164713-A 25 07-SEP-2001; Gasche, Christoph (AT); Zakeri, Schaker M. (AT) Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                  synthetic construct.
synthetic construct
artificial sequences.
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Sequence 25 from Patent WOO164713.
AX338660
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /gene="VHJ558-D:J"
/note="possible frameshift at D-J junction during rearrangement may result in nonfunctional immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                heavy chain"
9 c
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<1. .>50
                                                                                                                                                                                          /organism="synthetic construct"
/db_xref="taxon:32630"
/note="cDNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Mus musculus"
/strain="MRL +/+"
/db_xref="taxon:10090"
/clone="MPJ-09"
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/dev_stage="newborn"
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Post-processing: Minimum Match 0%
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Copyright (c) 1993 - 2003 Compugen Ltd.
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18	22	20	24	24.	24	<u>1</u> 3	24	24	19	19	24	24	24	24	1.8	24	19	19	22	22	21	21	21	20	22	22	21	22	21	21	22	16	16	22	24
AAT94899	æ	22	ABK67218	ABK41287	ABK40865	AAQ24360	ABL42407	32	AAV54819	AAV33786	AAI69204	ABI87281	ABI87280	ABK49469	AAT88795	ABQ77609	AAV50842	AAV50841	AAF67684	AAS03144	AAC89290	AAC88374	AAC87155	AAX88249	AAI62905	AAK65537	AAZ69526	AAH27041	AAA07136	AAA88764	AAH76860	AAT08198	AAT08201	10	ABL92859
Hu	Hu	Hu	Hu	Hu	Hu	2	Hu	M30	PCR	In	Hu	Са	Ca	пн	Fe	Hu	Br	Br	· Ch	Hu	Pr	Pr	Ri	Hu	Hu	Hu	Hu	. In	s.	Re	Hu	Am	Amp	PCI	o o
genomic	single nuc	biallelic p	qene specif	USF2 gene b	SS	ide pr	an alpha	protein ve	primer of	in beta-1 c	Human tyrosinase 1	Capture oligonucle	Capture oligonucle	Human cell divisio	Feline parvoviral	lutama	asp.	Brassica sp. polym	n insulator	novel	G1.	r G1.	EPSPS P		Human genomic DNA		Ð	terleukin	pyogenes	н	Human tyrosinase 1	lification	plification	R primer flt1D2.	protein-coupled

## ALIGNMENTS

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RESULT 1
ABA05479/c
                                                                                                                                                                          Human; RNA gyrase 12; malignant tumour; haemopathy; HIV; infection; human immunodeficiency virus; immunological disease; inflammation; enzyme; PCR primer; ss.
                                                                                                                                                                                                                                   01-MAR-2002
                                                                                 02-MAR-2000; 2000CN-0111862.
                                                                                                  02-MAR-2000; 2000CN-0111862
                                                                                                                    05-SEP-2001.
                                                                                                                                       CN1311325-A
                                                                                                                                                                                                               Human RNA gyrase 12 PCR primer SEQ ID NO 4.
                                                                                                                                                                                                                                                     ABA05479;
                                                                                                                                                                                                                                                                       ABA05479 standard;
                                                                                                                                                        Homo sapiens.
                                                              (BODE-) BODE GENE DEV CO LTD SHANGHAI
                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                       DNA;
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RESULT 2
ABN89564/c
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Matches 18
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Best Local S
Matches 19
                                                                                                                                             The present invention describes human uracil mononucleotide synthetase 9.9 (I). Also described is a process for producing (I) using DNA recombination technology. (I) and the polynucleotide encoding it can be used in the treatment of various diseases. The present sequence represents a PCR primer for (I), which is used in an example from the
                                                                                                                                                                                                                                     A human uracil mononucleotide synthetase 9.9 polypeptide, polynucleotide encoding it, for treating various diseases
                                                                                                               Sequence 24
                                                                                                                                                                                                                                                                                                                                           07-JUL-2000; 2000CN-0117079
                                                                                                                                                                                                                                                                                                                                                                                                            CN1333341-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human uracil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABN89564
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABN89564 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 24
                                                                                                                                                                                                                                                                          WPI; 2002-340719/38
                                                                                                                                                                                                                                                                                                                                                                07-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; uracil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        this polypeptide and DNA recombinant techniques to produce this polypeptide. The present invention also discloses a method of applying this polypeptide to treat various diseases, such as malignant tumour, haemopathy, HIV infection, immunological diseases and various
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example
                                                                                                                                     present invention.
                                                                                                                                                                                                                   Example
                                                                                                                                                                                                                                                                                                                                                                                       30-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                     (SHAN-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             inflammations. The present sequence is that of a PCR primer, useful to
                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention.
                         23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24
                                                                     18;
                                                                                                                                                                                                                 2; Page 16 (Disclosure); 34pp; Chinese.
                                                                                                                                                                                                                                                                                                Xie Y;
                                                                                                                                                                                                                                                                                                                      SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2; Page 17
                         CCTGGGCAAGAGTGAGACTCC
                                     CCTGGGTAAGAGTGGGACACC 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATCCTGGGTAAGAGTGGGACACC 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AGCCTGGGTGAGAGTGAGACCCC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 3 A; 10 C; 6 G; 5 T; 0 other;
                                                                                                               BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                  2000CN-0117079
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mononucleotide synthetase 9.9 PCR primer 2 SEQ ID NO:4
                                                                                                                                                                                                                                                                                                                                                                                                                                                       mononucleotide synthetase
                                                                                                                                                                                                                                                                                                                      BIODOOR GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
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                                                                                                               9 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            55.3%;
82.6%;
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8 5
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                                                                                                               6 G;
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                                                                                                               5 T;
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                                                                              16.2;
No. 6.
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                                                                                                               0 other;
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                                                                6.5e+02;
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RESULT 4
AAF89248
ID AAF8
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AAF89248 standard;

DNA;

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ВP

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RESULT 3
AAZ66299
В
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                                                                Query Match
Best Local
                                                   Matches
                                                                                                                                                                                               invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human genome; biallelic marker; high density disequipgenomic map; haplotype; phenotype; polymorphic base; haplotyping; hybridisation; identification; character diagnosis; single nucleotide polymorphism; SNP; ds.
                                                                                                        Sequence
                                                                                                                                                                                     effects from pharmaceutical agents acting on a disease as well as other
                                                                                                                                                                                                                                                                                                                                                                 Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                        map of the human genome -
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cohen D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-APR-1998;
23-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human map-related biallelic marker SEQ ID NO:646
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ66299;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAZ66299 standard;
                                                                                                                                            and 3367, are not actually given
                                                                                                                                                                                                                                                                                                                                   AAZ65654 to AAZ69578 represent human biallelic markers from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                          treatment
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                 2000-013267/01
 10
                                                  l Similarity
19; Conser
               CATCCTGGGTAAGAGTGGGACACC
                                                                                                        47
                                                                                                                                 present invention
                                                                                                                                                            SEQ ID NOS 2852,
                                                                                                                                                                                                                                                                                                                                                               Page 368;
CATCCTGGCTAACAGAGTGAAACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Blumenfeld
                                                                                                        BP;
                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     biallelic marker; high density disequilibrium map;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99WO-IB00822
                                                                                                        16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98US-0109732
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /standard_name= "single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            replace(24,C)
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                                                                                                        A; 14 C;
                                                                                                                                                                                                                                                                                                                                                              2745pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
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                                                              53.3%;
79.2%;
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                                                                                                                                                             2913,
                                                                                                        8 G;
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                                                                 Pred.
                                                                            Score
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n a sequence
                                                Mismatches
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 33
                                                                 No.
                                                                            16;
                                                                                                        0 other;
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                                                  3.9e+02;
5;
                                                                            DB 21;
                                                                                                                                              3096, 3
in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              characterisation; SNP; ds.
                                                                                                                                                                                                                                                                                                                                                                                                     high density disequilibrium
                                                                           Length 47;
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                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polymorphism"
                                                                                                                                              Sequence Listing
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RESULT 5
AAT25713
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Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes methods of clustering members of a sample, involving applying a hierarchical clustering algorithm to the sample members, determining the optimal number of clusters based on this and distributing the sample members into clusters using non-hierarchical clustering: The methods are useful in population based studies such as clinical trials, DNA fingerprinting and genetic profile analyses. The present samples was most to describe the sample members into clustering and several profile analyses.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Genetic clustering by distributing members into optimal numbers clusters determined by a hierarchical clustering algorithm or by paired pair analysis of homozygous pairs in clusters got from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
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07-JUL-2000; 2000US-0216897.
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  Homo
                                                                   Gene signature; messenger RNA; mRNA; relative abundance; frequency; human; cloning; mapping; non-biased library; diagnosis; detection;
                                                                                                                                                                                                  10-OCT-1996
                                                                                                                                                                                                                                                                                                              AAT25713 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
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non-hierarchical
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                                                                                                                                                    Human gene signature HUMGS07919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-OCT-2000; 2000WO-IB01632
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present sequence was used to demonstrate the method of the invention.
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sapiens
                                             typing;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3 CATCCTGGGTAAGAGTGGGACACC 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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gle nucleotide polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                              CATCCTGGCTAACAGAGTGAAACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page 71; 100pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 16 A; 14 C; 8 G; 9 T; 0 other;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     γď
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of
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RESULT 6
AAD31162/c
ID AAD311
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A single-stranded DNA (or its complementary strand or the corresp. Couble-stranded DNA) which comprises one of the 7837 "GS" sequences given in AAT19001-T26837 and which is able to hybridise to part of human genomic DNA, CDNA or mRNA is claimed. The GS (Gene Signature) sequences were obtained from 3'-directed CDNA ibraries prepared from various human tissues; synthesis of cDNA was initiated from the 3'-end of mRNA by using poly(T) as the sole primer. Since the 3'-en untranslated sequence is unique to a particular mRNA, species, almost all the 3'-oriented cDNAs hybridise with specific mRNAs. Each library is constructed so as to reflect accurately the relative abundance of different mRNAs in the particular tissue from which it was derived. The appearance frequency of a given GS in a cDNA library can be determined (esp. using primers and probes derived from the GS sequences) as a means of diagnosing abnormal cell function or for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                             episodic and paroxysmal anxiety disorder; Parkinson's disease; multiple sclerosis; Alzheimer's disease; cardiovascular disease; heart failure; angina pectoris; kidney disease; dyslipidaemias; obesity; emesis; gastrointestinal disorder; inflammatory bowel disease; diabetes; osteoporosis; inflammation; infection; human immunodeficiency virus; HIV cancer; inmune disorder; urinary retention; asthma; allergy; arthritis; benign prostatic hypertrophy; endotoxin shock; sepsis; gene therapy;
                               gynaecological disorder;
                                                                                                                                                                psychiatric disorder; central nervous system disorder; schizophrenia;
                                                                                                                                                                                    colon;
                                                                                                                                                                                                                                 Primer IP14924 for cloning human IGS43 GPCR cDNA
                                                                                                                                                                                                                                                                                                                                       AAD31162 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 1912; 2245pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               for diagnosis of abnormal cell function, by preparing cDNA that reflects relative abundance of corresp. mRNA in specific human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Identifying gene signatures in 3'-directed human cDNA library - for diagnosis of abnormal cell function, by preparing cDNA that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         W09514772-A1
                                                                                                                                                                                                                                                                      27 - AUG - 2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1995-206931/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matsubara
                                                                                                                                                                                                  IGS43; G-protein coupled receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MATS/) MATSUBARA K.
(OKUB/) OKUBO K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                       17 TCCAGNCTGGGTAACAGTGCGACTC
                                                                                                                                                                                    small
                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 TTCATCCTGGGTAAGAGTGGGACAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ζ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          different cell types
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                              intestine; stomach; mammary gland; prostate;
                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Okubo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     93JP-0355504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      94WO-JP01916
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12 A; 13 C;
                                                                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      53.3%;
76.0%;
                                                                                                                                                                                                                                                                                                                                        29
                               vaccine; human; PCR primer;
                                                                                                                                                                                                                                                                                                                                       ΒP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12 G; 12 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                  GPCR; uterus; lung;
                                                                                                                                                                                                                                                                                                                                                                                                                         41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         œ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                  trachea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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WO200228897-A2

obesity;

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                                                                                                                                                                                                                                                                                   dyslipidaemias, obesity, emesis, gastrointestinal disorders including irritable bowel syndrome, inflammatory bowel disease, diabetic gastroparesis and diabetes, ulcers, diarrhoea, osteoporosis, inflammations, infections including bacterial, fungal, protozoan and viral infections, particularly human immunodeficiency virus (HIV)-1 or HIV-2 infections, pain, cancers, chemotherapy induced injury, tumour invasion, immune disorders, urinary retention, asthma, allergies, complications of diabetes mellitus, and gynaecological disorders. The polypeptide and polynucleotide of the invention are also useful as research reagents and materials for discovery of treatments and diagnostics to animal and human diseases. The polynucleotide is also useful for Chromosome identification. The polypeptide is also usef
                                                                                                                                                                                                                              for assessing the binding of small molecule substrates and ligands in cells, cell-free preparations, chemical libraries and natural product mixtures. The present sequence is a PCR primer used for cloning human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      disorder, epilepsy, migraine, attention deficit/hyperactivity disorder, cardiovascular diseases including heart failure, angina pectoris, arrhythmias, myocardial infarction, cardiac hypertrophy, hypotension, hypertension, thrombosis, arteriosclerosis; cerebral vasospasm, subarachnoid haemorrhage, cerebral ischaemia, cerebral infarction,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            dyskinesias, Tourette's syndrome, tics, tremor, dystonia, spasms, anorexia, bulimia, stroke, addiction/dependency/craving, sleep
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel G-protein coupled receptor, termed IGS43 polypeptide and nucleic acid encoding the polypeptide, useful for treating disorders of uterus kidney, lung, colon, stomach, mammary gland, prostate and testis -
                                                                                                                                                                   Sequence 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               anxiety disorder, autism, delirium, multiple sclerosis, Alzheimer's disease, dementia, severe mental retardation, Huntington's disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           schizophrenia, episodic and paroxysmal anxiety disorders e.g. obsessive compulsive disorder, post traumatic disorder, phobia and panic, major depressive disorder, bipolar disorder, Parkinson's disease, general
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              intestine, stomach, mammary gland, prostate, testis, central nervous system, cerebellum and spinal cord. The polypeptide is useful as vaccine for inducing immunological response in a mammal, for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 33; 59pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-OCT-2000; 2000EP-0203411
04-OCT-2000; 2000US-237394P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             psychiatric and central nervous system disorders including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 polynucleotide encoding it. The IGS43 polypeptide, polynucleotide, agonist, antagonist or antibody is useful for treating dysfunctions or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to IGS43 G-protein coupled
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Deleersnijder W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    peripheral vascular disease, Raynaud's disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (SOLV ) SOLVAY PHARM BV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-SEP-2001; 2001WO-EP11319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         isorders related to uterus, kidney, lung, trachea, colon,
                                                                                                                                                                                                               GPCR cDNA
27
                                                                                                    Similarity
                                       AGAGTGGGACACCTGTG
                                                                                                                                                                   BP; 5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Blockx H,
                                                                                                  51.3%;
94.1%;
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                                                                                                    Score 15.4; DB 24
Pred. No. 1.6e+03;
                                                                                                                                                                   æ
                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Moor
                                                                                                                                                                   T; 0 other;
                                                                                                                        DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    kidney disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             receptor (GPCR) and the
                                                                                Indels
                                                                                                                      Length
                                                                                                                          29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  small
                                                                                                                                                                                                                                                                                              is also useful
                                                                                0;
                                                                            Gaps
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RESULT 7

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                                                                                                                                                                                                                                                                        dyskinesias, Tourette's syndrome, tics, tremor, dystonia, spasms, anorexia, bulimia, stroke, addiction/dependency/craving, sleep disorder, epilepsy, migraine, attention deficit/hyperactivity disorder, cardiovascular diseases including heart failure, angina pectoris, arrhythmias, myccardial infarction, cardiac hypertrophy, hypotension, arrhythmias, myccardial infarction, cardiac hypertrophy, hypotension,
                                                                   irritable bowel syndrome, inflammatory bowel disease, diabetic gastroparesis and diabetes, ulcers, diarrhoea, osteoporosis, inflammations, infections including bacterial, fungal, protozoan and viral infections, particularly human immunodeficiency virus (HIV)-1 or
                                                                                                                                                                                                                                                                                                                                                                                                                                                               for inducing immunological response in a mammal, for treating psychiatric and central nervous system disorders including schizophrenia, episodic and paroxysmal anxiety disorders e.g. obses compulsive disorder, post traumatic disorder, phobia and panic, maj depressive disorder, bipolar disorder, Parkinson's disease, general
                HIV-2 infections, pain, can invasion, immune disorders,
                                                                                                                                                                            dyslipidaemias, obesity, emesis, gastrointestinal disorders including
                                                                                                                                                                                                                               hypertension, thrombosis, arteriosclerosis, cerebral vasospasm, subarachnoid haemorrhage, cerebral ischaemia, cerebral infarcti
                                                                                                                                                                                                                                                                                                                                                                                                               anxiety disorder, autism, delirium, multiple sclerosis, Alzheimer's disease, dementia, severe mental retardation, Huntington's disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              intestine, stomach, mammary gland, prostate, testis, central nervous system, cerebellum and spinal cord. The polypeptide is useful as vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel G-protein coupled receptor, termed IGS43 polypeptide and nucleic acid encoding the polypeptide, useful for treating disorders of uterus, kidney, lung, colon, stomach, mammary gland, prostate and testis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-426102/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-OCT-2000; 2000EP-0203411.
04-OCT-2000; 2000US-237394P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         osteoporosis; inflammation; infection; human immunodeficiency virus; cancer; immune disorder; urinary retention; asthma; allergy; arthrit; benign prostatic hypertrophy; endotoxin shock; sepsis; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     epioodic and paroxysmal anxiety disorder: Parkinson's disease; multiple sclerosis; Alzheimer's disease; cardiovascular disease; heart fallure; anglna pectoris; kidney disease; dyslipidaemias; obesity; emesis; gastrointestinal disorder; inflammatory bowel disease; diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polynucleotide encoding it. The IGS43 polypeptide, polynucleotide, agonist, antagonist or antibody is useful for treating dysfunctions or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Deleersnijder W, Blockx
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           colon; small intestine; stomach; mammary gland; prostate; testis;
psychiatric disorder; central nervous system disorder; schizophrenia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAD31164 standard; DNA;
                                                                                                                                                                                                       peripheral vascular disease, Raynaud's disease, kidney disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disorders related to uterus, kidney, lung, trachea, colon,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to IGS43 G-protein coupled receptor (GPCR) and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-SEP-2001; 2001WO-EP11319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gynaecological disorder; vaccine; human; PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human IGS43 GPCR DNA amplifying primer IP15332.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (SOLV ) SOLVAY PHARM BV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IGS43; G-protein coupled receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      G-protein coupled receptor; GPCR; uterus; small intestine; stomach; mammary gland; p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2; Page 36; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                           pain, cancers, chemotherapy induced injury, tumour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       De Moor
                                                                                                                                                                                                                               infarction,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 trachea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Small
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        obsessive
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major

QΥ

Query Match Best Local

Sequence 29 BP; 4 A;

IGS43 GPCR DNA.

Matches

16;

Conservative

insecticide

gene

Similarity

В

.27

AAA91086/c

AAA91086 standard;

DNA;

Flt1

leakage;

WO200075319,-A1 Unidentified plasma 05-APR-2001 AAA91086; RESULT 8

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Qγ
                   Matches
                         Query Match
Best Local 9
3 CATCCTGGGTAAGAGTGGGACACCTGTG 30
                         Similarity
                         50.7%;
                 0;
                 Score 15.2; D
Pred. No. 2e+0
0; Mismatches
                         2e+03;
                 8
                 Indels
                 0,;
                 Gaps
```

ABL92859 standard;

.DNA;

22

ВP

45;

0;

Gaps

0;

Sequence 38

permeability

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complications of diabetes mellitus, and gynaecological disorders. The polypeptide and polynucleotide of the invention are also useful as research reagents and materials for discovery of treatments and diagnostics to animal and human diseases. The polynucleotide is also useful for chromosome identification. The polypeptide is also useful for assessing the binding of small molecule substrates and ligands in cells, cell-free preparations, chemical libraries and natural product mixtures. The present sequence is a PCR primer used for amplifying human
                                                                                                                                                    sequences encoding the fusion proteins of the invention between the Fill receptor and the Fc region of IgG. The specification relates to modified chimeric polypeptides with improved pharmacokinetics. The modified chimeric polypeptides are preferably Fill receptor polypeptides that have been modified to improve their pharmacokinetic profile. The polypeptides can be used to decrease or inhibit plasma leakage and/or
                                                                                                                                                                                                                                                                               This sequence represents a PCR primer used to construct DNA
                                                                                                                                                                                                                                                                                                                    Example 17; Page 63; 159pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acid molecule encoding mammalian phospholipid transfer protein, and its fragments, useful for diagnosis, evaluation, and treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-MAY-2000; 2000WO-US14142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR primer Flt1D2.VEGFR3D3.s for Flt1 receptor fusion DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                 diseases associated with the gene expression and for producing model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (REGE-) REGENERON PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14 AGAGTGGGACACCTGTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AGAGAGGGACACCTGTG
                                                                                                  BP; 9 A; 9 C; 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         fusion protein; chimeric; vascular permeability; I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-0138133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            51.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10 C; 8
                                                                                                                                         sed to decrease in a mammal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Yancopoulos
                                                                                                  G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 15.4;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       grotein; pharm
IgG Fc region;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .6e+03;
                                                            DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB
                                                          22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          pharmacokinetic;
                                                        Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR
                                                              38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 9
AAA28244/c
ID AAA282
ABL92859
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        B
                                                                                                  DЪ
                                                                                                                                       VО
                                                                                                                                                                                                                                                                                                                        PI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PD
XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PAX
                                                                                                                                                                                                  Query Match
Best Local
                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                pseudohyphal growth or hyphal growth. The invention further comprises determining whether the protein encoded by the essential gene has homology to a mammalian, plant or insect protein. The methods are useful for identifying genes in Saccharomyces cerevisiae which are essential for germination and proliferation of S. cerevisiae and using the identified genes or their encoded proteins as targets for highly specific antifungal agents, insecticides, herbicides and antiproliferation drugs. The formation includes examples of the use of the method, comprising the formation and characterisation of a number of S. cerevisiae genes to determine if they are essential for growth of the yeast. The present sequence represents a PCR primer used in the construction and analysis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gene useful for the design or discovery of an antifungal agent. The method comprises disrupting the function of a gene in a yeast cell, and identifying whether the function is essential for germination, growth,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Identifying a target gene for design or discovery of an antifungal agent, insecticide, or herbicide, comprising disrupting the function a gene in a yeast cell and identifying whether the function is essential for e.g. germination -
                                                                                                                                                                                                                                                            Sequence 45 BP; 20 A; 10 C; 7 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention relates to a method for the identification of a target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Saccharomyces cerevisiae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Upstream45 primer used in the construction of mutant YJR012C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA28244;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA28244 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200058457-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Essential gene identification; antifungal agent;
                                                                                                                                                                                                                                                                                                    the S. cerevisiae YJR012C mutant strain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ROSE-) ROSETTA INPHARMATICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     growth; pseudohyphal growth;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nerbicide;
                                                                                                  45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29
                                                                                                                                                                                 17;
                                                                                                                                         3 CATCCTGGGTAAGAGTGGGA
                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTTCCTGGGCAACAGCTGGATATCTATG
                                                                                                  CATTCTTGGTAAGAGTCGGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer;
                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2000WO-US08641
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99US-0127272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             156pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
                                                                                                                                                                                                  50.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  rowth; hyphal growth; YJR012C; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВP
                                                                                                                                                                             0;
                                                                                                                                                                                                  Score 15.2; DB
Pred. No. 2e+03;
                                                                                                  26
                                                                                                                                         22
                                                                                                                                                                                 Mismatches
                                                                                                                                                                                             No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gent; yeast; germination;
proliferation; insectici
                                                                                                                                                                                                                    21;
                                                                                                                                                                                                                    Length
                                                                                                                                                                               Indels
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tunction

08-JUN-1999;

WPI; 2001-071076/08

Ľ,

Davis

ABL92859

06-JUN-2002

(first

entry)

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05-JAN-2001

05-JAN-2001

08-JAN-2001

09-JAN-2001

10-JAN-2001

10-JAN-2001

22-JAN-2001

23-JAN-2001

23-JAN-2001

23-JAN-2001

20-FEB-2001
                           disease, especially cancer associated with altered levels of GPCR proteins and polynucleotides, by measuring the level of protein expression or the amount of nucleic acid from a mammal and compari with another mammal not having or not predisposed to the disease.
                                                                                         GPCR associated disorders such as cardiomyopathy, atherosclerosis, or a disorder related to cell signal processing and metabolic pathway modulation, in humans. GPCR proteins and the polynucleotides encoding them are useful for determining the presence of or predisposition to a
                                                                                                                                                               and antiinflammatory. They can be used in gene therapy and vaccine production. The GPCR proteins can be used for treating or preventing
                                                                                                                                                                                  (GPCR) designated GPCR1-36 from the present invention. The GPCRs can have activities such as: antidiabetic; anorectic; immunomodulator; cytostatic; neuroprotective; nootropic; antiparkinsonian; analgesic; osteopathic; immunosuppressive; metabolic; ophthalmological; antibacterial; virucide; fungicide; protozoacide; hypertensive; hypotensive; anti-HTV; antiulcer; antiasthmatic; antilipaemic; antiallergic; antiinfertility; haemostatic; antianfertility; haemostatic; antianfertility; haemostatic; antianfertility; haemostatic; antianfertility; haemostatic; and antiinflammatory. They can be used in gene therapy and vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Spytek KA,
Casman SJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-AUG-2000;
11-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              റ
proteins
                                                                                                                                                                                                                                                                                                                                                                             New G-protein coupled receptor polypeptides and nucleic acids, useful for diagnosis, prevention or treatment of hematopoietic, neurodegenerative, immune and signal transduction pathway disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               immunosuppressive; ophthalmological; antibacterial; virucide; fungicide; protozoacide; hypertensive; hypotensive; analgesic; osteopathic; antiulcer; antiasthmatic; antiallergic; anti-HIV; antilipaemic; vaccine; antiinfertility; antinflammatory; haemostatic; cell signal processing; cardiomyopathy; atherosclerosis; metabolic pathway modulation; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200212343-A2
                                                                                                                                                                                                                                                                                                                                              Example 2; Page 357; 492pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Larochelle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CURA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunomodulator; neuroprotective; nootropic; antiparkinsonian;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein-coupled receptor GPCR10 PCR primer
                                                                                                                                                                                                                                                                                                             present invention describes novel human
                                                                                                                                                                                                                                                                                                                                                                                                                                                       2002-217180/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          G protein-coupled receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURAGEN CORP
          are also useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ΨJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Padigaru M,
Vernet CAM,
Mezes PS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2001US-260283P
2001US-260450P
2001US-261156P
2001US-263334P
2001US-263434P
2001US-263704P
2001US-269964P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2001WO-US24787
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2001US-287916P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2001US-274873P
2001US-276406P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2001US-260072P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2001US-260003P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000US-224815P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000US-224613P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Taupier
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    prımer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Zerhusen BD, Baumgartner JC, Li
Ballinger RA, Shenoy SG, Kekuda
Grosse WM, Alsobrook JP, Gorman I
RJ, Colman SD, Szekeres ES;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SS
          identifying an agent,
                           or not predisposed to the disease. GPCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antidiabetic;
                                                                                                                                                                                                                                                                                                          G protein-coupled receptors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          anorectic;
          especially cellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NO:341
                                           and comparing it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     72 F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cytostatic;
an; metabolic;
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В

N

TCCTGGGCAACAGCTGGATATCTATG

27

RESULT 12

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RESULT 11
AAA91083
Qy
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                        Matches
                                              Query Match
                                     Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                sequences encoding the fusion proteins of the invention between the Flt1 receptor and the Fc region of IgG. The specification relates to modified chimeric polypeptides with improved pharmacokinetics. The modified chimeric polypeptides are preferably Flt1 receptor polypeptides that have been modified to improve their pharmacokinetic profile. The polypeptides can be used to decrease or inhibit plasma leakage and over vascular permeability in a mammal.
                                                                      Sequence 38 BP;
                                                                                                                                                                                                                                     Nucleic acid molecule encoding mammalian phospholipid transfer and its fragments, useful for diagnosis, evaluation, and treatm diseases associated with the gene expression and for producing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               candidate substance interacting with an olfactory receptor polypeptic its fragments or variants. The present sequence represents a PCR prinused in the isolation of a novel human GPCR in the present invention
                                                                                                                                                                                                  Example 17; Page 62; 159pp; English
                                                                                                                                                                                                                            systems
                                                                                                                                                                                                                                                                                     WPI; 2001-071076/08
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             plasma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Flt1 receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAA91083 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 22 BP; 7 A; 5 C; 6 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                             Papadopoulos NJ,
                                                                                                                                                                                                                                                                                                                                    (REGE-)
                                                                                                                                                                                                                                                                                                                                                                                  23-MAY-2000; 2000WO-US14142
                                                                                                                                                                                                                                                                                                                                                                                                          14-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200075319-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA91083
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    receptor or a downstream effector that binds to GPCR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
 u
                        19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             leakage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ū
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3 CATCCTGGGTAAGAGTGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16;
                                                                                                                                                                                                                                                                                                                                    REGENERON PHARM INC
                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
TCCTGGGTAAGAGTGGGACACCTGTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CATCCTGGGTAAGAAAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Flt1D2.VEGFR3D3.as
                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fusion protein;
; vascular perme;
                                                                                                                                                                                                                                                                                                                                                           99US-0138133
                                                                      8 A; 6 C; 10 G; 14 T; 0 other;
                                                                                                                                                                                                                                                                                                             Davis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
                                     49.3%;
73.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         49.3%;
88.9%;
                                                                                                                                                                                                                                                                                                           s,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             permeability;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВΡ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20
                        0,:
                                                                                                                                                                                                                                                                                                           Yancopoulos GD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
                                     Pred. No.
                                              Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for Flt1 receptor fusion DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chimeric protein; pharmacokinetic;
ability; IgG Fc region; PCR primer;
                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                14.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14.8;
 30
                                     3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.8e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB
                                              DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                   and treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    for screening
                                                38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           polypeptide,
a PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence
                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                       model
                                                                                                                                                                                                                                                            protein,
                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SS
                       0
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AAT08201/c
ID AAT08201 standard; DNA; 38
XX
AC AAT08201;
XX
AC AAT08201;
XX
CART08201;
XX
Linker; interleukin-4; in
KW granulocyte-colony stimul;
KW cancer; therapy; granulocyte
polymerase chain reaction
XX
PD 19-OCT-1995.
XX
O6-APR-1994; 94US-02252
XX
PPR (USSH) US DEPT HEALTH &
XX
O8-APR-1994; 94US-02252
XX
PPR (USSH) US DEPT HEALTH &
XX
O8-APR-1995 PR
XX
Circularly permuted ligar
PT interleukin(s) or colony-
PT Circularly permuted ligar
PT interleukin(s) or colony-
PT binding specificity and a
XX
AT08176-T08219 represent
CC AAT08176-T08219 represent
CC another joined to a lini
CC interleukin-4 (II-4), IL
CC (G-CSF) and granulocyte a
CC sequence joined to a lini
CC sequence was used in con
CC of IL-2 for use in a mod
CC another protein (preferr
CC sequence was used in con
CC of IL-2 for use in a mod
CC another protein (preferr
CC sequence was used in con
CC of IL-2 for use in a mod
CC another protein (preferr
CC sequence was used in con
CC of L-2 for use in a mod
CC another protein (preferr
CC sequence was used in con
CC of L-2 for use in a mod
CC where the two copies of
XX
Sequence 39 BP; 7 A; 10
                                                                                                                                                                                           RESULT 13
AAT08198
      XXX DXX AX ID
                                                                                                                                                                                                                                                                             B
                                                                                                                                                                                                                                                                                                                  δÃ
                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (G-CSF) and granulocyte macrophage-colony stimulating factor (GM-CSF). The linker regions used are represented by AR87726 and AAR87730. This sequence was used in conjunction with AAT88200 to amplify codons 39-127 of IL-2 for use in a modified ligand. These modified ligands are known as circularly permuted ligands. The modified ligands can be joined to another protein (preferrably a cytotoxin or an antibody) using a spacer (see AAR87732-R87734). The circularly permuted ligands have greater specificity and binding affinity than the original ligand. They can be used in chimeric molecules to inhibit the growth of tumour cells in vivo, where the two copies of the protein sequence are joined to cytotoxins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Circularly permuted ligands and chimeric mols. are modified interleukin(s) or colony-stimulating factors - with increa binding specificity and affinity for inhibiting tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Linker; interleukin-4; interleukin-2; IL-4; IL-2; G-CSF; GM-CSF; granulocyte-colony stimulating factor; circularly permuted ligand; cancer; therapy; granulocyte_macrophage-colony stimulating factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT08176-T08219 represent amplification primers used in the construction of modified ligands. The modified ligands comprise two copies of a sequence joined to a linker. The sequences used in the ligand include
                                                                                                                                                                       AAT08198 standard; DNA;
Linker; interleukin-4; interleukin-2; IL-4; IL-2; G-CSF; GM-CSF;
                                         Amplification primer BK135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequence joined to a linker. The sequences used in the ligand in interleukin-4 (IL-4), IL-2, granulocyte-colony stimulating factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 5; Page 35; 97pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (USSH ) US DEPT HEALTH & HUMAN SERVICES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Amplification primer BK138
                                                                                       28-MAY-1996
                                                                                                                                                                                                                                                                           26
                                                                                                                                                                                                                                                                                                                  2 TCATCCTGGGTAAGAGTGGGACACCT 27
                                                                                                                                                                                                                                                                                                                                                              . Similarity
19; Conserv
                                                                                                                                                                                                                                                                           TCACCGGAGGTAACGGTGGGGCACCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            94US-0225224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95WO-US04468
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       reaction;
                                                                                                                                                                                                                                                                                                                                                                               49.3%;
73.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                            10 C; 12 G; 10 T; 0 other;
                                                                                                                                                                       42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English.
                                                                                                                                                                       ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR; primer; amplify; ss.
                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                               Score 14.8;
Pred. No. 36
                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                               3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                 DB 16;
                                                                                                                                                                                                                                                                                                                                                          7;
                                                                                                                                                                                                                                                                                                                                                                                                 Length
                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               increased
                                                                                                                                                                                                                                                                                                                                                                                                   39;
                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             construction
                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                          0
  Вþ
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AAH76860/c
                                                                                                                                                                                                                                                                                                                                          RESULT 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 19
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WO200171001-A1
                                     Homo sapiens.
                                                                                         cytostatic; anti-HIV;
                                                                                                                               Human; tyrosinase 15; recombinant production;
malignant tumour; cancer; blood disease; HIV infection;
                                                                                                                                                                                                                           14-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 5; Page 35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Circularly permuted ligands and chimeric mols, are modified interleukin(s) or colony-stimulating factors - with increased binding specificity and affinity for inhibiting tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               W09527732-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      granulocyte-colony stimulating factor; circularly permuted cancer; therapy; granulocyte macrophage-colony stimulating polymerase chain reaction; PCR; primer; amplify; ss.
                                                                                         human immunodeficiency virus; immune disorder; inflammatory
cytostatic; anti-HIV; antiinflammatory; immunomodulator;
                                                                                                                             malignant tumour;
                                                                                                                                                                                     Human
                                                                                                                                                                                                                                                                                                   AAH76860 standard; DNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1995-366354/47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kreitman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (USSH ) US DEPT HEALTH & HUMAN SERVICES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-OCT-1995
                                                                                                                                                                                     tyrosinase
                                                                                                                                                                                                                                                                                                                                                                                                 Ģ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         l Similarity
19; Conser
                                                                        transcription-PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RJ,
                                                                                                                                                                                                                                                                                                                                                                                               TCACCGGAGGTAACGGTGGGGCACCT
                                                                                                                                                                                                                                                                                                                                                                                                                                  TCATCCTGGGTAAGAGTGGGACACCT 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 10 A; 12 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94US-0225224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         95WO-US04468
                                                                                                                                                                                   15 RT-PCR primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   97pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         49.3%;
73.1%;
                                                                                                                                                                                                                                                                                                     ВP
                                                                          RT-PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14.8; DB 16;
Pred. No. 3.1e+03;
0; Mismatches 7;
                                                                        primer;
                                                                                                                                                                                     SEQ
                                                                                                                                                                                     IJ
                                                                          SS
                                                                                                                                                                                     NO:4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ligand;
                                                                                                             condition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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RESULT 15
AAA88764
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to human tyrosinase 15 (AAG66766), nucleic acids encoding it (AAH76858), and a method for the recombinant production of tyrosinase 15. The protein has a molecular weight of 15 kb. The present invention additionally discloses an antagonist of tyrosinase 15 for therapeutic use, and an antibody which specifically binds to tyrosinase 15. Tyrosinase 15, and nucleotides which encode it may be used for treating a variety of diseases, such as malignant tumours, blood diseases, HIV (human immunodeficiency virus) infection, immune disorders and inflammatory conditions. The protein may also be used to screen for modulators of its activity or for peptide fingerprinting identification. The polynucleotide can be used as a primer for nucleic acid amplification reactions or as a probe for hybridisation reactions, or in producing gene chips or microarrays. Sequences AAH76859-AAH7680 represent reverse transcription-PCR (RT-PCR) primers used in an exemplification of the
                                                                                                                                                                                                                                             Key
5'UTR
19-OCT-2000.
                                                                                                                                                                        CDS
                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-FEB-2001
                                                WO200061611-A2
                                                                                                                                                                                                                                                                                                                                                                                                                   RSV; vaccine; attenuation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Respiratory syncytial virus NS1-NS2 gene fusion region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA88764 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human tyrosinase 15 and encoded polynucleotide, used in diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and inflammation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Мао
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                                                                                                                                                                                                                                                                                                                                         Human respiratory syncytial virus
                                                                                                                                                                                                                                                                                                                                                                                            NS2 gene; mutant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-MAR-2000; 2000CN-0115044
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (BIOW-) BIOWINDOW GENE DEV INC SHANGHAI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        invention to isolate human tyrosinase 15 cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6 CCTGGGTAAGAGTGGGACACC 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CCTGGGTGACAGTGCGAGACC 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Page 11; 33pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 3 A; 9 C; 8 G; 4 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2001WO-CN00357
                                                                                              /partial
/note= "
                                                                                                                                         /*tag=
                                                                                                                                                                                          note= "NS1 nontranslated region
                                                                                                                                                                                                                      /*tag=
                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                            SS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              48.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   \mathfrak{S}
                                                                                         end of NS2 open reading
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                                                                                                                                                                                                 pneumonia; bronchiolitis; NS1 gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 14.6;
Pred. No. 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                              frame'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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Вþ

9

TCCTTGGTTAGAGATGGACAC

29

0

Search completed: June 21, Job time : 167.531 secs

2003, 22:26:18

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Qy
                                                                                                                                                                                                  Matches
                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents a fusion of human respiratory syncytial virus (RSV) NS1 gene 5' nontranslated region to the NS2 coding region. PCR-mediated mutagenesis (see AAA88760-61) was used to delete the NS1 gene coding region, 3' untranslated region, the NS1-NS2 coding region and the NS2 5' nontranslated region, a cotal of 529 nucleotides. The deleted region corresponds to nucleotides 99-627 of the RSV antigeneme cDNA given in AAA88743. This type of deletion can be incorporated into viable recombinant RSV of the invention to yield an altered phenotype, in this case reduced rate of virus growth and reduced plaque size in vitro. The invention relates to infectious chimeric RSV produced by introducing 1 or more heterologous gene(s) or gene segment(s) from 1 RSV subgroup or strain into a recipient RSV brookground of a different subgroup or strain into a recipient RSV produced by a different subgroup or strain into a recipient RSV produced by a different subgroup or strain into a recipient RSV produced by a different subgroup or strain into a recipient RSV produced by a different subgroup or strain into a recipient RSV produced by a different subgroup or strain into a recipient RSV produced by a different subgroup or strain into a recipient RSV produced by a different subgroup or strain into a recipient RSV produced by a different subgroup or strain into a recipient RSV produced by a different subgroup or strain into a produced by a different subgroup or strain into a produced produced by a different subgroup or strain into a produced produced by a different subgroup or strain into a produced produced by a different subgroup or strain into a produced produced by a different subgroup or strain into a produced produced by a different subgroup or strain into a produced produced by a different subgroup or strain and subgroup or strain into a produced produced by a different subgroup or strain and subgroup or subgroup or str
                                                                                                                                                                                                                                                                                             particle of select
                                                                                                                                                                                               Sequence 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Infectious chimeric respiratory syncytial virus (RSV) produced from cloned nucleotide sequences, useful as a vaccine against diseases caused by the virus, such as pneumonia and bronchiolitis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                        subgroup or strain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 17; Fig 21; 280pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-679462/66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Collins PL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (USSH)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-APR-1999;
                                                                                                                                                                                                                                                                                    bgroup or strain. The resulting chimeric RSV virus or subviral rticle is infectious and attenuated, preferably by introduction selected mutations. The chimeric RSV is useful as a vaccine
         տ
                                                                    l Similarity
17; Conserv
                                                                                                                                                                                                                                                          RSV, which causes pneumonia and bronchiolitis in infants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US DEPT HEALTH & HUMAN SERVICES
      TCCTGGGTAAGAGTGGGACAC 25
                                                                                                                                                                                               BP;
                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Murphy BR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2000WO-US08802
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-0291894
                                                                                                                                                                                               12 A; 7 C; 6 G; 8 T; 0 other;
                                                                                              48.7%;
81.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Whitehead SS;
                                                                    Score 14.6; [Pred. No. 3.7e 0; Mismatches
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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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1: /cgn2_6/ptodata/1,

2: /cgn2_6/ptodata/1,

3: /cgn2_6/ptodata/1,

4: /cgn2_6/ptodata/1,

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length: 50
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/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
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                                              US-08-454-557C-63
US-08-340-426D-63
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US-08-25-24-31
US-08-25-24-28
US-08-25-24-28
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US-08-25-24-28
US-08-31-22-38
PCT-US95-04468-28
PCT-US95-0459-3-38
PCT-US95-0599-3-12
US-09-71-425-7
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TTCATCCTGGGTAAGAGTGGGACACCT	imilarity ; Conserva	ON FOR SEQ ID NO: E CHARACTERISTICS: H: 31 base pairs nucleic acid DEDNESS: both OGY: both 7C-63	AGENT Ludwig ATION CE/DOO NICATION	TYPE: F R: IBM I RC SYSTEM E: Patem PPLICATION NUM TION NUM DATE: 33	Washington D.C. U.S.A.	WALL NVENT SEQUI	-63 Ap			4444	444444
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Best Local Similarity
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APPLICANT: de la Monte,
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        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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APPLICANT: Wands, Jack R.
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
                                                                                                                                                                                                                                            TITLE OF INVENTION:
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APPLICANT: Wands, Jack R.
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74.18;
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Pred. No. 93;
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PCT-US95-17111A-63/c
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Best Local Similarity
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                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 31 base pairs
TYPE: nucleic acid
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APPLICATION NUMBER: 08/
FILING DATE: 14-NOV-199
                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: PCT/US95/17111A.
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MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS:
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LENGTH: 31 base pairs
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                                                                                                                                                                                                                   NAME: Ludwig, Steven R. REGISTRATION NUMBER: 36, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                  FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                           STRANDEDNESS:
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Conservative
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74.1%;
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74.1%;
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0;
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              Score 15.8;
Pred. No. 9:
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Mismatches
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Gaps
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                                                                                                                                                                                                                                                                            RESULT 6
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; MOLECULE TYPE:
US-08-225-224-31
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                                                                                                                                                                                                Patent No. 6011002
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Best Local Similarity
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                                                   NERAL INFORMATION.
APPLICANT: Pastan, Ira
APPLICANT: Kreitman, Robert J.
APPLICANT: Kreitman, Robert J.
APPLICANT: Kreitman, Robert J.
APPLICANT: Puti, Raj K.
TITLE OF INVENTION: Circularly Permuted Ligands and
TITLE OF INVENTION: Circularly Permuted Chimeric Molecules
TITLE OF SECUENCES: 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 39 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen L.
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 15:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPUTER:
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: PASTAN, ITA
APPLICANT: KREITMAN, R
TITLE OF INVENTION: CI
TITLE OF INVENTION: CI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/
FILING DATE: 8-APR-1994
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o. 6011002
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5. 5635599
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                 San Francisco
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California
                                E: Townsend and Townsend and Crew LLP Two Embarcadero Center, Eighth Floor
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Steuart Street Tower, One Market Plaza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            : (415) 543-96
(415) 543-5043
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CIRCULARLY
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73.1%;
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Pred. No. 2
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Matches
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           REFERENCE/DOCKET NUMBER: 15
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 543-9600
                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US (
FILING DATE: 08-APR-1994
                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
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ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
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PRIOR APPLICATION DATA:
PRIOR APPLICATION NIMBER: WO PCT/US95/04468
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MEDIUM TYPE: Floppy disk
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE:
                                                                NAME: Weber, Ellen L. REGISTRATION NUMBER:
                                                                                                                                                                                              APPLICATION NUMBER: PCT/US95/04468 FILING DATE: 07-APR-1995
                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                               CLASSIFICATION:
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OTHER INFORMATION:
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73.18;
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RESULT 9
US-08-722-258-28
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Sequence 28; Application US/08722258 Patent No. 6011002
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Best Local
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Best Local Similarity
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SEQUENCE CHARACTERISTICS:
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NAME: Weber, Ellen L.
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 155
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/
FILING DATE: 8-APR-1994
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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MEDIUM TYPE: Floppy disk
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TOPOLOGY: linear
NT.ECULE TYPE: DNA (primer)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
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T: Steuart Street Tower, One Market Plaza
San Francisco
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5635599
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EDNESS: single
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CIRCULARLY PERMUTATED LIGANDS AND
CIRCULARLY PERMUTED FUSION PROTEINS
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73.1%;
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Pred. No. 2
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Pred. No. 2.9e+02;
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                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                               TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
               SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 01
FILING DATE: 08-APR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION DATA:
PRIOR APPLICATION UNIMBER: WO PCT/US95/04468
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 72
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Circularly Permuted Ligands and TITLE OF INVENTION: Circularly Permuted Chimeric Molecules
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                             LOCATION: 1..42 OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: WO PILLING DATE: 06-APR-1995
APPLICATION NUMBER:
                               SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
TOPOLOGY: li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME:
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                                                                                                                                                                                                                                                                                                                                                            l Similarity
19; Conserv
                                                                                                                                                                                                                                                                                                                          2 TCATCCTGGGTAAGAGTGGGACACCT 27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Weber, Ellen Lauver
                                                                                                                                                                                                                                                                                          TCACCGGAGGTAACGGTGGGGCACCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
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Puri, Raj K.
                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pastan, Ira
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IBM PC compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 single
                                                                                                                                 CIRCULARLY PERMUTATED LIGANDS AND CIRCULARLY PERMUTED FUSION PROTEINS
                                                                                                                                                                                                                                                                                                                                                                            49.3%;
73.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "BK-135 primer"
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PCT/US95/04468
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Pred. No. 2.
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                               Version #1.30
                                                                                                                                                                                                                                                                                                                                                                            .9e+02
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07-APR-1995

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 38, Application  
Patent No. 6030835
GENERAL INFORMATION:
                TELEFAX: (415)
INFORMATION FOR SEQ ID NO: ?
SEQUENCE CHARACTERISTICS:
SEQUENCE TENGTH: 45 base pairs
1.FNGTH: 45 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local :
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 42 base pairs
TYPE: nucleic acid
                                                                                                  REFERENCE/DOCKET NUMBER: BA
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 926-6200
                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Musser M.D., James M.
APPLICANT: Kapur M.D., Vivek
TITLE OF INVENTION: Methods and Compositions for Identifying
TITLE OF INVENTION: Group A Streptococcus
                                                                                                                                                                                         APPLICATION NUMBER: US 08/160,965 FILING DATE: 02-DEC-1993 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 14-SEP-1994
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 543-9600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE:
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                      NAME: Rae-Venter Ph.D., Barbara REGISTRATION NUMBER: 32,750
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Menlo Park
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Weber, Ellen L. REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 94025-7022
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CLASSIFICATION:
STRANDEDNESS:
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o. 6030835
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     E: WEIL, GOTSHAL & MANGES 2882 Sand Hill Road, Suite 280
                                                                                     (415) 854-3713
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     USA
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                                                                                                                                                                                                                                                                  US 08/306,542
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                                                                                                                                       BAYL-004/01US
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Pred. No. 2.9e+02;
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US-08-931-220-38
                                                                        ; IMMEDIATE SOURCE: ; CLONE: speB19 PCT-US95-11723-38
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Query Match
Best Local Similarity
Thehes 20; Conserve
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; Sequence 38, Application PC/TUS9511723
; GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
4E PRICE PAIRS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                  ANTI-SENSE: NO ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                         FILING DATE: 14-SEP-1994
ATTORNEY/AGENT INFORMATION:
NAME: Rac-Venter Ph.D., Barbara
REGISTRATION NUMBER: 32,750
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Streptococcus pyogenes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORIGINAL SOURCE:
                                                                                                                                                              HYPOTHETICAL:
                                                                                                                                                                             MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: BAYL-004/03US TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Musser M.D., APPLICANT: Kapur M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HYPOTHETICAL: NO ANTI-SENSE: NO
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                                                                                                                     ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: FILING DATE: 14-SEF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE:
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CITY: Palo Alto
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nes 20; Conservative
                                                                                                                                                                                                          STRANDEDNESS:
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                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                                TELEPHONE:
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                                                                                                                                                                                                                                        LENGTH:
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                                                                                                                                                                                                                                                                                  (415) 424-8760
               Conservative
                                                                                                                  Streptococcus pyogenes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PatentIn Release #1.0, Version #1.30
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14-SEP-1995
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                                                                                                                                                                             DNA (genomic)
                                                                                                                                                                                                       single
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69.0%;
                           48.7%;
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Identifying Streptococcus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   James M.
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Pred. No. 3.7e+02;
                           Score 14.6; DB 5;
Pred. No. 3.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
               Mismatches
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                                         Length 45;
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               Indels
             0;
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2 TCATCCTGGGTAAGAGTGGGACACCTGTG 30

TCATGTTGGGGAAGGGCGGGAACACTTTG 5

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RESULT 14
US-09-742-703-12
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                                                                                                                                                           Query Match
Best Local S
Matches 20
Sequence 12, Application US/09742703 Patent No. 6423543
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                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (415) 854-37
NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 45 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
ZIP: 94025-7022
COMPUTER READABLE FORM:
COMPUTEN TYPE: Floppy disk
                                                                                                                                                                                                                                                     IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                    ORIGINAL SOURCE:
ORGANISM: Streptococcus pyogenes
                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: BA
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 926-6200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/306,542
FILING DATE: 14-SEP-1994
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 0 FILING DATE: 02-DEC-1993 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                     HYPOTHETICAL:
ANTI-SENSE: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
                                                                                                                                                            Local Similarity
les 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: FILING DATE: 01-MA
                                                                                                                                                                                                                                                                                                                                                     ropology:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                      CULE TYPE:
                                                                                              33 TCATGTTGGGGAAGGGCGGGAACACTTTG
                                                                                                                           2 TCATCCTGGGTAAGAGTGGGACACCTGTG 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rae-Venter Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Application PC/TUS9605997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: WEIL, GOTSHAL & MANGES 2882 Sand Hill Road, Suite 280
                                                                                                                                                                                                                                                                                                                                                                                                                                                (415) 854-3713
                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fernandez,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kapur M.D., Vivek
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Musser M.D., James M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ananthaswamy, H. N.
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01-MAY-1995
                                                                                                                                                                                                                                                                                                                                     DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                    single
                                                                                                                                                                          48.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            us 08/160,965
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32,750
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                                                                                                                                                                          Score 14.6;
Pred. No. 3.
                                                                                                                                                            Mismatches
                                                                                                                                                                        .7e+02;
                                                                                                                                                                                            DB 5;
                                                                                                                                                                                        Length 45;
                                                                                                                                                         Indels
                                                                                                                                                           0;
                                                                                                                                                         Gaps
                                                                                                                                                         0;
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NUMBER OF SEQ ID NOS: 49
SEQ ID NO 12
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
Search completed: June 22, 2003, 00:01:29 Job time: 38.625 secs
                                                                                              QΥ
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                                                                  В
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                                                                                                                                                                                             ; OTHER INFORMATION: Synthetically generated oligonucleotides US-09-171-425-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 15
US-09-171-425-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; OTHER INFORMATION: Antisense Oligonucleotide US-09-742-703-12
                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
LENGTH: 42
                                                                                                                               Matches
                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 2, Application US/09171425A Patent No. 6465438
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Best Local Similarity
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                                                                                                                                                Best
                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/171,425A CURRENT EILING DATE: 1998-10-19 EARLIER APPLICATION NUMBER: PCT/EP97/01943 EARLIER FILING DATE: 1996-04-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/742,703 CURRENT FILING DATE: 2000-12-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Patrick Allen | APPLICANT: Lex M. Cowsert TITLE OF INVENTION: ANTIS: FILE REFERENCE: RTS-0090
                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: NUCLEIC ACID VACCINATION FOR PARVOVIRAL INFECTIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 08909/003001
                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                           ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                            TYPE: DNA
                                                                                                                                             Local Similarity
                                                                14
                                                                                                                               19;
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                                                                                              2 TCATCCTGGGTAAGAGTGGGACACCTG 28
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                                                                                                                                                                                                                                                                                                                                                                                                                                Schorr, Joachim
Baker, Henry J.
Smith, Bruce F.
                                                               TCAACCTGGCTAAGAAGAAGAAAGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TCCTGGGCCAGGGTGGGAC 20
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70.4%;
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84.2%;
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq
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Perfect score:
                                                                                                                                                                                                                                                                                                                         Database :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        length: 0
length: 50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 TTCATCCTGGGTAAGAGTGGGACACCTGTG
                                                                                                                                                                                                                                                                      Published_Applications_NA:*

i: /cgn2_6/ptodata/2/pubpna/US07_PUBCOMB.seq:*

cgn2_6/ptodata/2/pubpna/PCT_NEW_PUB.seq:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compus
/cgin2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*
/cgin2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*
/cgin2_6/ptodata/2/pubpna/US09_NEW_PUB.seq:*
/cgin2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*
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/cgin2_6/ptodata/2/pubpna/US10_PUBCOMB.seq:*
/cgin2_6/ptodata/2/pubpna/US10_PUBCOMB.seq:*
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/cgn2_6/ptodata/2/pubpna/US07_NEW_PUB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.	, , , , , , , , , , , , , , , , , , ,	Score	% Query Match Length	Length	DB	ID	Description
	1	30	100.0	30	9	US-09-964-667-9	Sequence 9, Appli
	8	30	100.0	30	10	US-09-964-666-9	Sequence 9, Appli
	w	30	100.0	30	10	US-09-964-412-9	Sequence 9, Appli
O	4	22	73.3	22	9	US-09-964-667-5	Sequence 5, Appli
O	U	22	73.3	22	10	US-09-964-666-5	Sequence 5, Appli
c	0	22	73.3	22	10	US-09-964-412-5	Sequence 5, Appli
	7	15.6	52.0	25	ဖ	US-10-098-263B-65996	Sequence 65996, A
C	æ	15.2	50.7	45	10	US-09-965-602-27	Sequence 27, Appl
ဂ	9	15	50.0	43	ဖ	US-09-802-640-9	Sequence 9, Appli
	10	14.8	49.3	30	9	US-10-085-906-97	Sequence 97, Appl
	11	14.8	49.3	31	10	US-09-801-274-8	Sequence 8, Appli
	12	14.6	48.7	25	ဖ	US-10-215-112-7585	Sequence 7585, Ap
G	ω,	14.6	48.7	25	9	US-10-215-112-12152	Sequence 12152, A
0	14	14.6	48.7	25	ø	US-10-098-263B-97774	Sequence 97774, A
a	G	14.6	48.7	25	ø	US-10-098-263B-129375	Sequence 129375,
	16	14.4	48.0	49	9	US-09-860-670-233	Sequence 233, App
O	17	14.2	47.3	25	φ	US-10-098-263B-3274	Sequence 3274, Ap
C	818	14.2	47.3	36	ø	US-10-294-778-8	Sequence 8, Appli
	19	14.2	47.3	39	9	US-10-011-672-9	Sequence 9, Appli

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13.2	13.2	13.2	13.4	13.4	13.4	13.4	13.4	13.4	13.4	13.4	13.4	13.6	13.6	13.6	13.6	13.6	13.6	13.6	13.6	13.8	14	14	14	14.2	14.2
44.0	44.0	44.0	44.7	44.7	44.7	44.7	44.7	44.7	44.7	44.7		45.3	45.3	45.3	45.3	45.3	45.3	45.3	45.3	46.0	46.7	46.7	46.7	47.3	47.3
25	25	25	40	40	40	31	30	25	25	25	25	25	25	25	25	25	25	25	25	31	31	25	25	39	39
9	9	Ø	10	10	φ	9	9	9	9	9	9	9	9	9	9	9	9	9	9	10	9	9	9	9	9
US-10-098-263B-19558	US-10-060-895A-1149	US-10-060-895A-1148	US-09-922-101-60	US-09-951-401-60	US-09-951-402-60	US-09-864-636A-2477	US-10-082-772-9	US-10-098-263B-125737	US-10-098-263B-103293	US-10-098-263B-26737	US-10-098-263B-26559	US-10-098-263B-103720	US-10-098-263B-31788	US-10-060-895A-1147	US-10-060-895A-1146	US-10-060-895A-1145	US-10-060-895A-1144	US-10-060-895A-1143	US-10-060-895A-1142	US-09-801-274-873	US-09-912-263-86	US-10-098-263B-65995	US-10-098-263B-44679	US-10-012-070A-9	US-10-012-013-9
 19558,	Sequence 1149, Ap	Sequence 1148, Ap	Sequence 60, Appl	Sequence 60, Appl	Sequence 60, Appl	Sequence 2477, Ap	Sequence 9, Appli	Sequence 125737,	Sequence 103293,	Sequence 26737, A	Sequence 26559, A		31788,	1147, A	6, A	Sequence 1145, Ap	1144, A	Sequence 1143, Ap	1142, 1	Sequence 873, App	ā.		Sequence 44679, A	Sequence 9, Appli	Sequence 9, Appli

## ALIGNMENTS

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US-09-64-667-9
Sequence 9, Application US/09964667
Publication No. US20030033621A1
GENERAL INFORMATION:
                                                                                                                                                                          INFORMATION FOR SEQ ID NO: 9:
                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 2005-3934

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/964,667
                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 0609.4370000
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
TELEPAX: 202-371-2540
                                                                                                                                                                                                                                                                                                                                                FILING DATE: 28-Sep-2001 CLASSIFICATION: <Unknown> ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox, P.L.L.C.
STREET: 1100 New York Ave., Suite 600
                            TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Transgenic Animals and Cell Lines for
Screening Drugs Effective for the Treatment or Preventio
of Alzheimer's Disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: de la Monte, Suzanne
Wands, Jack R.
SEQUENCE DESCRIPTION: SEQ ID NO: 9:
                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                 NAME: Esmond, Robert W. REGISTRATION NUMBER: 32,893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: DC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
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of Alzheimer's Disease

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RESULT 3
US-09-964-412-9
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                                                                     Sequence 9, Application US/09964412
Patent No. US20020129391A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 9, Application US/09964666 Patent No. US20020104108A1 GENERAL INFORMATION:
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TELEFAX: 202-371-2540
INFORMATION FOR SEQ ID NO: 9:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox, P.L.L.C. STREET: 1100 New York Ave., Suite 600
TITLE OF INVENTION: Transgenic Animals and Cell Lines for Screening Drugs Effective for the Treatment or Prevention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Transgenic Animals and Cell Lines for Screening Drugs Effective for the Treatment or Prevention
                                     APPLICANT: de la Monte, Suzanne
Wands, Jack R.
                                                                                                                                                                                                                                                                                                                                                     SEQUENCE DESCRIPTION: SEQ ID NO: 9:
                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/09/964,666
FILING DATE: 28-Sep-2001
CLASSIFFICATION: <UDknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: de la Monte, S
Wands, Jack R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30;
                                                                                                                                                                                                               1 TTCATCCTGGGTAAGAGTGGGACACCTGTG 30
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                                                                                                                                                                                               TTCATCCTGGGTAAGAGTGGGACACCTGTG 30
                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Esmond, Robert W.
REGISTRATION NUMBER: 32,893
REFERENCE/DOCKET NUMBER: 06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Washington
                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                 Score 30; DB 10;
Pred. No. 0.00015;
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Pred. No. 0.00015;
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RESULT 4
US-09-964-667-5/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 5, Application US/09964667 Publication No. US20030033621A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION
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COMPUTER READABLE FORM:
COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
APPLICATION NUMBER: US/09/964,667
FILING DATE: 28-Sep-2001
CLASSIFICATION: duknown>
ATTORNEY/AGENT INFORMATION:
                                                                            SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS: ADDRESSEE: Sterne,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: 202-371-2600 TELEFAX: 202-371-2540
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ADDRESSEE: Sterne, Kessler,
STREET: 1100 New York Ave.,
                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: de la Monte, Suzanne
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 TTCATCCTGGGTAAGAGTGGGACACCTGTG 30
                                                                                                                                                                                                                                                            CITY: Washington
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REGISTRATION NUMBER: 32,893
REFERENCE/DOCKET NUMBER: 06
                                                                                                                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                    ZIP: 20005-3934
                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                             STATE: DC
                                                                                                                                                                                                                                                                                   ADDRESSEE: Sterne, Kessler, Goldstein & Fox, P.L.L.C
STREET: 1100 New York Ave., Suite 600
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ilarity 100.0%;
Conservative
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                                                                                                                                                                                                                           USA
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Pred. No. 0.00015;
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                                                                                                     Version #1.30
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; TOPOLOGY: linear; MOLECULE TYPE: CDNA SEQUENCE DESCRIPTION: SEQ ID NO: 5: US-09-964-667-5
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US-09-964-666-5/c
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                                  Matches
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Best Local S
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Best Local Similarity
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                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 5: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: de la Monte, Suzanne
Wands, Jack R.
TITLE OF INVENTION: Transgenic Animals and Cell Lines
Screening Drugs Effective for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox, P.L.L.C STREET: 1100 New York Ave., Suite 600
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REFERENCE/DOCKET NUMBER: 1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
                                                                                                                                                                                                                                                                NAME: ESMOND, ROBERT W.
REGISTRATION NUMBER: 32,893
REFERENCE/DOCKET NUMBER: 0609.4370000
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                   MOLECULE TYPE:
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                           cn 73.3%;
1 Similarity 100.0%;
22; Conservative (
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3 CATCCTGGGTAAGAGTGGGACA 24
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                                                                                                                                                                                 TYPE: nucleic acid
                                                                                                                                                                                                                                                    TELEFAX: 202-371-2540
                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/964,666 FILING DATE: 28-Sep-2001
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STRANDEDNESS: single
                                                                                                                                                                 STRANDEDNESS: single
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nilarity 100.0%;
Conservative (
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                               ; Score 22; DB
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0; Mismatches
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Pred. No.
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0.74;
                                                DB 10; Length 22; 0.74;
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; SEQUENCE DESCRIPTION: SEQ ID NO: US-09-964-412-5
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                                                                                                                                                                                                                                                                                           US-10-098-263B-65996
CURRENT APPLICATION NUMBER: US/10/098,263B
CURRENT FILING DATE: 2003-01-08
PRIOR APPLICATION NUMBER: 60/276,759
PRIOR FILING DATE: 2001-03-16
NUMBER OF SEQ ID NOS: 131066
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 65996
LENGTH: 25
                                                                                                                                                                                                                            Sequence 65996, Application US/10098263B Publication No. US20030104410A1 GENERAL INFORMATION:
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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                              APPLICANT: Mittman, Michael
TITLE OF INVENTION: Human Microarray
FILE REFERENCE: 3118.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 5: SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: US/09/964,412
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Floppy disk
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Wands, Jack R.
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STRANDEDNESS: single
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TELEFAX: 202-371-2540
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          73.3%;
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; Pred. No.
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US-10-098-263B-65996

TYPE: DNA
ORGANISM: Homo sapien

Query Match

DB 9;

Length 25

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                                                                                                                        US-09-802-640-9
                                                 Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                         CÜRRENT APPLICATION NUMBER: US/09/802/640
CÜRRENT FILING DATE: 2001-03-09
NÜMBER OF SEQ ID NOS: 122
SÖFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 9
                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 9, Application US/09802640 Publication No. US20030036057A1 GENERAL INFORMATION:
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PRIOR FILING DATE: EARLIER FILING DATE: 2000-03-31
NUMBER OF SEQ ID NOS: 68
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 27
LENCTH: 45
TYPE: DNA
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Matches
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Best Local S
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                                                                                                                                                                                                                                                                                                              APPLICANT: KLEYN PALTICK
TITLE OF INVENTION: GENES AND POLYMORPHISMS ASSOCIATED WITH
TITLE OF INVENTION: CARDIOVASCULAR DISEASE AND THEIR USE
FILE REFERENCE: 24736-2048
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Dimster-Deak, Dago
TITLE OF INVENTION: ESSENTIAL GENES IN YEAST AS TARGETS FOR ANTIFUNGAL
TITLE OF INVENTION: AGENTS,
TITLE OF INVENTION: HERBICIDES, INSECTICIDES AND ANTI-PROLIFERATION D
                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Braun, Andreas
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CURRENT FILING DATE: 2001-09-27
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                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                         LENGTH: 43
TYPE: DNA
ORGANISM: Artificial Sequence
PEATURE:
                                                                                                                                       OTHER INFORMATION: PCR primer
                                                 Local Similarity
mes 18; Conserv
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                8 TGGGTAAGAGTGGGACACCTGTG 30
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81.8%;
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                                             Score 15; DB y; LC. Pred. No. 1.3e+03; ....
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Pred. No. 6.9e+02;
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Pred. No. 1.
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                                                                                  Length 43;
                                                   Indels
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RESULT 12

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Вb
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: SEQ ID NO 8
: LENGTH: 31
: TYPE: DNA
: ORGANISM: Homo sapiens
US-09-801-274-8
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US-10-085-906-97
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Best Local S
Matches 19
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SEQ ID NO 97
LENGTH: 30
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                                                                             Query Match
Best Local Similarity
Matches 19; Conserv
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                                                                                                                                                                                                                                                                                                                   APPLICANT: Cargill, Michele
APPLICANT: Ireland, James S.
APPLICANT: Ireland, James S.
APPLICANT: Lander, Eric S.
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
FILE REFERENCE: 2825.2009-001
CURRENT APPLICATION NUMBER: US/09/801,274
CURRENT FILING DATE: 2001-03-07
PRIOR APPLICATION NUMBER: US/09/187,510
PRIOR APPLICATION NUMBER: US/60/206,129
PRIOR APPLICATION NUMBER: US/60/206,129
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TITLE OF INVENTION: POLYMORPHIC ELEMENTS IN THE
TITLE OF INVENTION: COSTINULATORY RECEPTOR LOCUS AND USES THEREOF
FILE REFERENCE: GRN-5343CP2
CURRENT APPLICATION NUMBER: US/10/085,906
CURRENT FILING DATE: 2002-02-27
                                                                                                                                                                                                                                                                               PRIOR FILING DATE: 2000-05-22
NUMBER OF SEQ ID NOS: 1802
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PRIOR FILING DATE: 1999-03-25
PRIOR APPLICATION NUMBER: US 09/534,061
PRIOR FILING DATE: 2000-03-24
PRIOR APPLICATION NUMBER: PCT/US00/07938
PRIOR FILING DATE: 2000-03-24
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APPLICANT: Wu, Pau
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les 19; Conserv
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CATCAAGGGATGAARTGTGACACCTGCG 29
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                                                                             Conservative
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73.1%;
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67.9%;
                                                                           Score 14.8; DB 10
Pred. No. 1.6e+03,
1; Mismatches
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Pred. No. 1.6e+03;
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                                                                                                                     DB 10;
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US-10-098-263B-97774/c: Sequence 97774, Application US/10098263B: Publication No. US20030104410A1; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                     RESULT 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: Synthetic Oligonucleotide US-10-215-112-12152
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Best Local S
Matches 17
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CURRENT APPLICATION NUMBER: US/10/215,112
CURRENT FILING DATE: 2002-08-08
NUMBER OF SEQ ID NOS: 14936
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 12152
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Best Local Similarity 81.0
Watches 17; Conservative
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Publication No. US20030082596A1
GENERAL INFORMATION:
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CURRENT FILING DATE: 2002-08-08
NUMBER OF SEQ ID NOS: 14936
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7585.
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                               APPLICANT: Mittman, Michael
TITLE OF INVENTION: Human Microarray
FILE REFERENCE: 3118.1
CURRENT APPLICATION NUMBER: US/10/098,263B
CURRENT FILING DATE: 2003-01-08
PRIOR APPLICATION NUMBER: 60/276,759
PRIOR FILING DATE: 2001-03-16
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                   NUMBER OF SEQ ID NOS: 131066
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ORGANISM: Artificial Sequence
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OFTWARE: Microarray Probe Sequence Listing Generator V 1.1
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81.0%;
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Pred. No. 2e+03;
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Query Match
Best Local Similarity
Thehes 17; Conserve
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Search completed: June 22, 2003, 03:18:14 Job time: 96.5938 secs
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US-10-098-263B-129375/c
                                                                                                                                                                                                                             SEQ ID NO 129375
LENGTH: 25
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                                                                                                                         Matches
                                                                                                                                     Query Match
Best Local :
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                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: U5/10/098,263B
CURRENT FILING DATE: 2003-01-08
PRIOR APPLICATION NUMBER: 60/276,759
PRIOR FILING DATE: 2001-03-16
                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Mittman, Michael TITLE OF INVENTION: Human Microarray FILE REFERENCE: 3118.1
                                                                                                                                                                                                                                                          SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 131066
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                                   M., Hultman, M., Kucaba, T., Rifkin, L., Rohlfing, T., Soan, R., Williamson, A., Wohldman The WashU-Merck EST Project Unpublished (1995)
Contact: Wilson RK
Washington University School
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                                                                          Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P. and Wilson,R.
                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 48)
                                                                                                                                         human.
                                                                                                                                                                       yp01g07.rl Soares breast 3NbHBst Homo sapiens cDNA clone IMAGE:186204 5' similar to gb|M87904|HUMALNES31 Human carcinoma cell-derived Alu RNA transcript, (rRNA); gb:M81695 LEUKOCYTE ADHESION GLYCOPROTEIN P150,95 ALPHA CHAIN (HUMAN);, mRNA sequence.
                       Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                   Homo sapiens
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                                    School of Medicine way, Box 8501, St. Louis,
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H84235 ys65f06 s1
AA976917 cq35h04|s
H14824 ym53d07 s1
AA410762 zt35e11|r
C20874 HUMGS000494
C20875 HUMGS000494
D12406 HUMOTW129
AX576739 AXT-27007
AA934082 cn27g11|s
AA928240 on87c07|s

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AR871520 vq36e07/r
H99413 yx24f02.\$1
T96362 ye49c04.\$1
AU104029 AU104029
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BI522018 603081749
D11989 HUM0005134
H39150 yn83e09.f1

linear

EST 16-AUG-1995

Hawkins,M., Holman M., Parsons,J.,

15

AU106864 AI439612

Email: est@watson.wustl.edu Insert Size: 730 High quality sequence starts: 1

MO 63108

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KEYWORDS
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                                                                FEATURES
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                                                                          Suzuki, Y., Yoshitomo-Nakagawa, K., Maruyama, K., Suyama, A., S. Construction and characterization of a full length-e a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156
                                                                                                                    Contact: Yutaka Suzuki
Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Jap
Email: Yuzuki@ims.u-tokyo.ac.jp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 TTCATCCTGGGTAAGAGTGGGACAC 25
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Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
Insert Length: 730
Std Error: 0.00
                                                                                                                                                                                                                       mapping of mRNA start sites
EMBO Rep. 2 (5), 388-393 (2001).
                                                                                                                                                                                                                                                                   Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J., Sese,J., Ha, Ota,T., ISogai,T., Tanaka,T., Morishita,S., Okubo,K., Sakaki,Y., Nakamura,Y., Suyama,A. and Sugano,S.
                                                                                                                                                                                                                                                     Diverse transcriptional initiation revealed by
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KAIA3269, mRNA sequence.
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/db_xref="taxon:9606"
                                                               Location/Qualifiers
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Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 50)
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AZ796512.1
                                                                       2M0052H18F Mouse 10kb plasmid UUGC1vclone UUGC2M0052H18 F, DNA sequence
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Jap
Email: ysuzuki@ims.u-tokyo.ac.jp
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house mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Yutaka Suzuki
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20; Conservative
                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                       /clone_lib="Sugano Homo sapiens cDNA library"
/note="Differential display comparison of untreated dimethylfumarate treated U937 cells"
10 c 11 g 22 t
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Sugano Homo sapiens
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/note="Differential display comparison of untreated and
dimethylfumarate treated U937 cells"
a 12 c 16 g 13 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone="COL03340"
                                      GI:12944647
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Best Local
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                                                                                         AU106864 Sugano Homo sapiens NBLAN21NR1, mRNA sequence.
                                                                                                                                                                                                                                                  2 TCATCCTGGGTAAGAGTGGGACACCTGTG 30
                                                                                                                                                                                                                                                                                                      21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Wellsr,R.
                                                        AU106864
AU106864.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           University of Utah Genome Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mammalia; Eutheria;
1 (bases 1 to 48)
 Homo sapiens
                                          EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     High quality sequence stop: 48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: ddunn@genetics.utah
Insert Length: 10000 std
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa;
                       numan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               was hydrodynamically sheared by repeated passage through a .005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gil4732114)gblaF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and the character of the complementary to the insert adaptors.
                                                                                                                                                                                                                                                                                                                                                                                                                   purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ambicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (http://www.jax.org/resources/documents/dnares/).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Laboratory Mouse DNA Resource
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//lab_host="E. Coli Strain XL10-Gold, T1-resistant, F
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/clone="UUGC2M0052H18"
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/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                              Score 16.2; DB 17;
Pred. No. 1.1e+04;
0; Mismatches 8;
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cDNA library Homo sapiens
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               High quality sequence stop:
Location/Qualifiers
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Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano,s. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
                           Trace considered overall poor quality Insert Length: 1378 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   tc91b01.x1 NCI_CGAP_CLL1 Homo similar to gb:M35531 GALACTOS:
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Suzuki,Y., Taira,H., Ts
,H., Ota,T., Isogai,T.,
Seq primer: -40UP from Gibco
                                                                                                                              M.D., Louis M. Staudt, M.D., Ph.D. CDNA Library Preparation: M. Bento Soares, Ph.D. CDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Clone distribution: NCI-GGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                        Tissue Procurement: Ash Alizadeh, John Byrd, M.D.,
                                                                                                                                                                                                                                                                                                                 Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                     Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                        www-bio.lln1.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                    Tumor Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                          NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
National Cancer Institute, Cancer Genome Ana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Yutaka Suzuki
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/note="Differential display comparison of untreated and
dimethylfumarate treated U937 cells"
a 14 c 15 g 11 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone="NBLAN21NR1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GI:4305543
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79.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GALACTOSIDE 2-L-FUCOSYLTRANSFERASE (HUMAN);,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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Pred. No. 1.4e+04;
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IMAGE:2073481 3'
                                                                                                                                                            information
                                                                                                                                                                                     Sequencing Center
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                                                                                                                                                                                                                                                                                              Mike
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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                                                                                                                                                                                                                                                                                              Grever,
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BASE COUNT
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Best Local Similarity
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Unpublished (1997)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 50)
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EST.
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                                                                                                                                                                                                                                                                                                                                                                                         CDNA Library Preparation: M. Bento Soares, Ph.D.
CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; 1st strand cDNA was prepared from pooled bulk breast tumor tissue, and was then primed with a Not I oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. This library is the normalized version of
NCI_CGAP_Brl.1.
                                                                                                                                                                     /clone_lib="NCI_CGAP_Br2"
/sex="female, pooled"
/tissue_type="breast"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                             /clone="IMAGE:1631407"
                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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/tissue_type="B-cell, chronic lymphotic leukemia"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone="IMAGE: 2073481"
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76.0%;
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Pred. No. 2.3e+04;
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Homo sapiens cDNA clone IMAGE:16314
repetitive element;, mRNA sequence
Library was constructed by Bento Soares
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VERSION KEYWORDS

ACCESSION DEFINITION

SOURCE

ORGANISM

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Matches

AI024893/c

RESULT 7

JOURNAL COMMENT

REFERENCE

AUTHORS TITLE

FEATURES

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50.0%; 72.0%;

Score 15; DB 14; Pred. No. 3.4e+04;

DB 14; Length 47;

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Query Match
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The WashU-Merck EST Project
Unpublisha
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              High qality sequence starts: I High qality sequence stops: 1 Source: IMAGE Consortium, LLNL This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Trace considered overall poor quality Insert Length: 716 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   yf18e09.s1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA cl. IMAGE:127240 3' similar to gb|m87933|HUMAALU364 Human carcinoma cell-derived alu RNA transcript, (rRNA);, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3 CATCCTGGGTAAGAGTGGGACACCT 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Washington University School of Medicine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (1995)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: est@watson.wustl.edu
Insert Size: 716
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                      and M. Fatima Bonaldo.
                                                                                                                                  (Pharmacia), digested with Pac I and cloned into the Pac I and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization Library
                                                                                                    constructed by Bento Soares and M. Fatima Bonaldo
                                                                                                                                                                                                                                                                                                                                      /db_xref="taxon:9606"
/clone="IMAGE:127240"
/clone_lib="Soares fetal liver spleen lNFLS"
                                                                                                                                                                                                                                                                                                                         /sex="male"
                                                                                                                                                                                                                                                                                                                                                                                            /db_xref="GDB:479401"
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Pred. No. 2.4e+04;
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                                                               Query Match
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N38850
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    High quality sequence stop: 1.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (1995)
Contact: Wilson RK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P. and Wilson,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IMAGE Consortium (info@image.llnl.gov
Trace considered overall poor quality
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
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314 286 1810
                                           Conservative
                                                                                                                                        double-stranded cDNA was size selected, ligated to Eco RI adapters (Pharmacia), digested with Not I and cloned into the Not I and cloned into the Not I and Eco RI sites of a modified pT7T3 vector (Pharmacia). Library went through one round of normalization to a Cot = 5. Library constructed by Bento Soares and M.Fatima Bonaldo. RNA from 4 multiple sclerosis lesions from one patient was kindly provided by Dr. Kevin G. Becker (NINDS/NIH)."
                                                                                                                                                                                                                                                                                                                                                           /tissue_type="multiple sclerosis lesions"
/dev_stage="Age 46"
/lab_host="DH1OB (empicillin resistant)"
/note="Vector: pT7T3D (Pharmacia) with a modified
polylinker V_TYPE: phagemid; Site_1: Not I; Site_2: E
: 1st strand CDNA was primed with a Not I - oligo(dT)
primer [5'
                                                                                                                                                                                                                                                                                                                                   /db_xref="GDB:3898260"
/db_xref="taxon:9606"
/clone="IMAGE:279884"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /sex="male"
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Pools of cells were isolated from a GenomeScreen(TM) library. The
library of cells was generated by retroviral integration of a gene
tagging element consisting of: 1) A promoterless beta-lactamase
proceeded by a splice acceptor as a reporter for gene expression;
2) A promoter driving neomycin resistance followed by a splice
donor to trap downstream exons. 3' RACE from neomycin gene was
performed using total RNA from isolated pools. Output was shotgun
cloned in pAmp-1 and used to transform DH5-alpha competent
bacteria. 5' ends of reported sequences were immediately preceded
                   Mus musculus
                                                                                                         AU258691\ 3'\mbox{-directed mouse cDNA} library Mus musculus BED0013497 3', mRNA sequence.
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, Durick,K. and Pollok,B
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1 (bases 1 to 40)
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AST-T32B0048 Genetrap T47D Human sapiens genomic 5', DNA sequence.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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Tel: 8584048436
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /cell_line="T47D"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /tissue_type="Carcinoma"
                                                                          GI:20324496
                                                                                                                                                                                                                                                                                                                                  49.3%;
73.1%;
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                                                                                                                                                                                                     Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A..and Sugano
,S. Construction and characterization of a full length-enriched and
a s'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
                                                                                                                                                                                                                                                                         Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Jaj
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AU108052 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone COLEGO15, mRNA sequence.
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Graduate School of Biological Sciences
Nara Institute of Science and Technology
8916-5 Takayama, Ikoma, Nara 630-0101, Japan
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Fax: 81-743-72-5589
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae
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URL:http://love2.aist-nara.ac.jp/BED/index.html.
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                                                                                                                                                                                                                                                                                                                                        Contact: Yutaka Suzuki
                                                                                                                                                                                                                                                                                                                                                            21270072
                                                                                                                                                                                                                                                                                                                                                                                                      Diverse transcriptional initiation revealed by fine, large-scale
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Generation of expressed sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                        Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J., Sese,J., Ha
,H., Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo,K., Sakaki
,Y., Nakamura,Y., Suyama,A. and Sugano,S.'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CATCCTGGCTAACACAGTGAAANCCGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and Matoba, R.
                                                      /clone_lib="Sugano Homo sapiens cDNA library"
/note="Differential display comparison of unt
dimethylfumarate treated U937 cells"
a 11 c 17 g 14 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /db_xref="taxon:1090"
/db_xref="taxon:1090"
/clone="BED0013497"
/clone_lib="3'-directed mouse cDNA library"
/tissue_type="brain"
/note="Vector: pGEM-T-easy"
15 c 7 g 10 t 1 others
                                                                                                                                               /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                /clone="COLE0015"
                                                                                                                                                                                                  Location/Qualifiers
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70.4%;
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            Score 14.8;
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Pred. No. 4.1e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tags
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            9
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          Length 50;
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                                                                                                                                                                                                                                                                                   Japan
                                                                                               of untreated
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          Query Match
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: est@watson.wusul.com
Trace considered overall poor quality
Possible reversed clone: similarity on wrong strand This clone is
Possible reversed clone: nvitrogen Corp. 2130 South Memorial
Tracker information call: (800)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Wylie,T., Underwood,K., Steptoe,M., Theising,B., Allen,M.,
,Y., Person,B., Swaller,T., Gibbons,M., Pape,D., Harvey,N.,
,R., Ritter,E., Kohn,S., Shin,T., Jackson,Y., Cardenas,M.,
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Glycine max
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 TTCATCCTGGGTAAGAGTGGGACACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              )-533-4363 or contact via email: ccu@resgen.com
Seq primer: -40RP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Washington University School of Medicine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PROTEIN. ;, mRNA sequence. AI900587
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Location/Qualifiers
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                                                                                                                                                                                              /note="Vector: pBluescript II XR; Site_1: EcoRI; Site_2: XhoI; This cDNA library was constructed from mRNA isolated from whole seedlings of 2-3 week old greenhouse grown plants. The cDNA library was prepared using the Stratagene pBluescript II XR cDNA library construction kit. Complementary DNA was synthesized from mRNA using a primer consisting of a poly (dT) sequence with a XhoI restriction site. EcoRI adapters were ligated to the blunt-ended cDNA fragments followed by XhoI digestion. The CDNA fragments were directionally cloned into the EcoRI-XhoI restriction was fixed to the control of the contr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               phyta; Magnoliophyta; eudicotyledons; core eudic
eurosids I; Fabales; Fabaceae; Papilionoideae;
Erpelding."
                                                                                                                        site of the pBluescript vector. The ligated cDNA fragments were transformed into XL10-Gold host cells. This library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                greenhouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone="GENOME SYSTEMS CLONE ID: Gm-c1013-513"
/clone_lib="Gm-c1013"
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/db_xref="taxon:3847"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 greenhouse grown"
/lab_host="XL10-Gold"
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    17
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9
                                                                                      Dr.
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L., Kucaba,T., Martin,J., Beck,C.
., Theising,B., Allen,M., Bowers
                                                                         Shoemaker and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Louis,
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48.78;

Score 14.6;

DB 9;

Length 40;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACCESSION
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                                                                       Matches
                                                                                          Best Local
                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 69.0 Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              JOURNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Trace considered overall poor quality Possible reversed clone: similarity on wrong strand Seq primer: -28ml3 rev2 ET from Amersham High quality sequence stop: 1.

Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL;
TMAGE Consortium (info@image.llnl.gov) for further i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Ld
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Marra,M., Hillier,L., Allen,M., Bowles,M., Dietrich,N., Dubu
Geisel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., Morris,M.,
Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B.,
Theising,B., Wylle,T., Lennon,G., Soares,B., Wilson,R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AA871520

46 bp mRNA linear EST 16-N vq36e07.rl Barstead bowel MPLRB9 Mus musculus cDNA clone IMAGE.1096356 5' similar to SW:GPR2_HUMAN P46092 PROBABLE GPROTEIN-COUPLED RECEPTOR GPR2. ; mRNA sequence.
                                                                      20; |Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MGI:602588
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (1996)
Contact: Marra M/Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Waterston,R.
The WashU-HHMI Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA871520.1 GI:2966965
                                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mammalia; Eutheria;
TTTACTCCGGGTACGAGGGGGGGGGGCCTAT 31
                                  TTCATCCTGGGTAAGAGTGGGACACCTGT 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TTCGTCCCCGTTCAGATCGGAACAGCTGT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TTCATCCTGGGTAAGAGTGGGACACCTGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                314 286 1800
314 286 1810
                                                                                                                                                                6
                                                                                                                                                                Barstead.
a 11 c
                                                                                                                                                                                                                                                                           3']; double-stranded cDNA was ligated to Eco RI adaptors [AATTCGGATCCTTG], digested with Not I and cloned into the Not I and Eco RI sites of the modified p77T3 vector. Source irradiated bowel harvested 72 hours after
                                                                                                                                                                                                                                                                                                                                                              /clone_lib="Barstead bowel MPLRB9"
/tissue_type="bowel"
/dev_stage="8 weeks"
/lab_host="DH10B"
                                                                                                                                                                                     irradiation (1400 Gys).
Barstead. "
                                                                                                                                                                                                                                                                                                                                                                                                                                        /db_xref="taxon:10090"
/clone="IMAGE:1096356"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /strain="EVB/N"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Mus musculus
                                                                                         48.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        69.0%;
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                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
                                                                                     Score 14.6; DB 9;
Pred. No. 4.9e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No. 4.6e+04; 0; Mismatches 9;
                                                                                                                                                               18
                                                                       Mismatches
                                                                                                                                                             ρ
                                                                                                                                                               11
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                                                                                                                                                                                                 Library constructed by
                                                                                                                                                               H
                                                                                                          9
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                                                                                                      Length 46;
                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     further information.
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                                                                      0,
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                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                      0,
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VERSION
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H99413
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMMENT
                                                                                                                                                                       ORIGIN
                                                                                                                                                                                        BASE COUNT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cocus
                                                                                                     Query Match
Best Local
                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             source
                                                                                   18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holma, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterst, R., Williamson, A., Wohldmann, P. and Wilson, R.

7. Williamson, A., Wohldmann, P. and Wilson, R.

7. WashU-Merck EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Source: IMAGE Consortium, LLNL;
This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further i
Trace considered overall poor quality
Insert Length: 679 Std Error: 0.00
                                         1 TTCATCCTGGGTAAGAGTGGGACAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 High quality sequence starts:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. Louis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          H99413 40 bp mRNA linear EST 15-DEC-1995 yx24f02.sl Soares melanocyte 2NbHM Homo sapiens cDNA clone IMAGE:262683 3' similar to gb:J02931 TISSUE FACTOR PRECURSOR (HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Insert Length: 679 Std Err
Seq primer: m13 -40 forward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (1995)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Н99413
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                                                                                                       Similarity
TCCAGCCTGGGNAAGAGAGCAAAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (bases 1 to 40)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   quality sequence stop: 1.
Location/Qualifiers
                                                                                                                                                                                          16
                                                                                   Conservative
                                                                                                                                                                                                                           double-stranded cDNA was size selected, ligated to Eco RI adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT7T3 vector (Pharmacia). Library constructed by Bento Soares and M.Fatima Bonaldo. RNA from normal foreskin melanocytes
                                                                                                                                                                                                                                                                                                                                (FS374) was kindly provided by Dr. Anthony P. Albino. 11 c 8 \ g \ 4 \ t \ 1 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone_lib="Soares melanocyte 2NbHM"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/db_xref="GDB:3872325"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone="IMAGE:262683"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /db_xref="taxon:9606"
                                                                                                  48.0%;
72.0%;
                                                                               0,
                                                                                                     Score 14.4;
Pred. No. 5
                                                                                   Mismatches
26
                                                                            5.5e+04;
7;
                                                                                                                      DB 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hominidae;
                                                                                                                        Length
                                                                                 Indels
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                                                                                                                        40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            contact the
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Search completed: June 21, Job time: 1302.03 secs